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Minimum DB :

length: 0 length: 2000000000

Potal number of hits satisfying chosen parameters:

1009251

1009251 seqs, 160854530 residues

Scoring table:

BLOSUM62 Gapop 10:0 , Gapext 0.5

Sequence: Perfect score: Title:

US-09-284-107-21 55

1 VLNASVTSPLLE 12

Run on: OM protein -

protein search, using sw model

Copyright (c) 1993 - 2000 Compugen Ltd.

June 4, 2001, 12:11:27; Search time 260.2 Seconds (without alignments) 7.418 Million cell updates/sec

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database

COMB. pep: \*

# ALIGNMENTS

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QY 1 VLNASVTSPLLE 12            Db 7 VLNASVTSPLLE 18	Query Match 100.0%; Score 55; DB 23; Length 96; Best Local Similarity 100.0%; Pred. No. 0.007; Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	; LOCATION: (1)(96) ; OTHER INFORMATION: Xaa = Any Amino Acid US-60-160-209-4252	ORGANISM: HUMAN FEATURE: NAME/KEY: VARIANT	; SEQ ID NO 4252 ; LENGTH: 96 ; TYPE: PRT	CURRENT APPLICATION NUMBER: US/60/160,209 CURRENT FILING DATE: 1999-10-19 NUMBER OF SEQ ID NOS: 4646 SOFTWARE: FastSEQ for Windows Version 4.0	0	RESULT 1 US-60-160-209-4252 ; Sequence 4252, Application US/60160209 ; GENERAL INFORMATION: ADDITION: BONAZZI VIVIEN	

Pred. No. is the number of results predicted by chance to rescore greater than or equal to the score of the result being and is derived by analysis of the total score distribution. number of results predicted by chance to have a an or equal to the score of the result being printed, SUMMARIES

/cgn2\_6/ptodata/2/paa/US60\_COMB.pep: /cgn2\_6/ptodata/2/paa/US097\_COMB.per

.6/ptodata/2/paa/US091\_COMB.pep:

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US-09-702-021-14 Sequence 14, A
                                                  GENERAL INFORMATION:
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Best Local Similarity
Matches 12; Conser
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                                                                                                                                                                                                                                                                                                                                            EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
   FILE REFERENCE:
                APPLICANT: C. Frase
                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Homo sapiens
-09-245-764-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 12; Conserv
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SOFTWARE: FastSEQ for
SEQ ID NO 4414
LENGTH: 102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/245,764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS FITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/60/160,203 CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Hogarth, P. Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ISOLATED HUM
TITLE OF INVENTION: NUCLEIC ACID
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                            ARLIER APPLICATION NUMBER: 60/073,972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICANT: McKenzie,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT: Powell,
                                                                                                                                                                                                                                                                                                                                    ENGTH: 261
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
                                                                                                                                            174 VLNASVTSPLLE 185
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/099,994 FILING DATE: 1998-09-11
                                                             Application US/09702021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maxwell, Kelly F. Garrett, Thomas P.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/09245764
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ilarity 100.0%;
Conservative (
                                   Fraser
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N: FAIL MOLECULES AND USES THEREOF 7853-217
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e, Ian F.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
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Pred. No. 0.022;
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                                                                                                                                                                                                      Mismatches
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US-09-107-532-4325
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                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY: SITE; LOCATION: (349); OTHER INFORMATION: PCT-US00-05882-1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Watches 12; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens US-09-702-021-14
                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn
SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 14
                                                                                                                                                equence 4325, Application US/09107532 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: PCT/US00/05882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Steve Ruben.
TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 37
                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (208)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: SITE
                                                          NUMBER OF SEQUENCES: 7
                                                                                                      FITLE OF INVENTION:
                                                                                                                                   APPLICANT:
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                                                                                        TITLE OF INVENTION:
                                STREET:
                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                            Local
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Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FastSEQ for Windows Version 3.0
                             100 Beaver Street
                                                                                            Lynn A Doucette-Stamm and David Bush
TENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING
TENTION: ENTEROCOCCUS FAECIUM FOR
                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ID NOS: 1694
                                         GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application
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                                                                        DIAGNOSTICS AND 7308
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                                                                                                                                                                                                                                                                                                                                                                                                                                  naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                      DB 1;
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URRENT APPLICATION DATA:

US/09/107,532

OPERATING SYSTEM

SOFTWARE:

COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660

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; LOCATION: 1...104
US-09-107-532-4325
                                                                         GRGANISM: Homo sapiens
JS-09-429-755-315
                                                                                                                                                                                                                                                                                                                                                                        JS-09-429-755-315
                                                                                                      SEQ ID NO 315
LENGTH: 292
TYPE: PRT
 Query Match 65.
Best Local Similarity 70.
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 65. Best Local Similarity 60. Matches 6; Conservative
                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/429,755A
CURRENT FILING DATE: 1999-10-28
NUMBER OF SEQ ID NOS: 315
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                     ITLE OF INVENTION:
                                                                                                                                                                                                           ILE REFERENCE:
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REFERENCE/DOCKET NUMBER: GTV
ELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RIOR APPLICATION DATA:
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FILING DATE: July 2, 1997
TTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                           INFORMATION
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                                                                                                                                                                                                                                                     Dillon, Davin
                                                                                                                                                                                                                                                                                                            Frudakis, Tony N. Smith, John M.
                                                                                                                                                                                                                                                                   Misher, Lynda
Retter, Marc W
                                                                                                                                                                                                                                                                                                Reed, Steven G.
                                                                                                                                                                                                                                                                                                                                                       Application US/09429755A
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                                                                                                                                                                                                           ON: TREATMENT AND DIAGNOSIS 210121.419C6
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                                                                                                                                                                                                                                      COMPOSITIONS AND METHODS FOR THE
               65.5%;
70.0%;
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 Score 36; DB Pred. No. 1.1e 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTC-012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
               1.1e+02;
                              DB 18;
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                                                                                                                                                                                                                        OF BREAST CANCER
                           Length 292;
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   Indels
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Gaps
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NUMBER OF SEQ ID NOS: 55
SOUTWARE: FastSEQ for Wi
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 70.0
Thes 7: Conservative
            NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ fo
SEQ ID NO 532
                                                                                                                                                                                                                                                                                                                                                                                     GENERAL
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                                                                 CURRENT APPLICATION NUMBER: US/09/483,672A CURRENT FILING DATE: 2000-01-14
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CURRENT FILING DATE: 1999-11-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE:
                                                                                                  FILE REFERENCE:
                                                                                                             PPLICANT: Meagher, Madeleine
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT
LENGTH: 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER TILE REFERENCE: 210121.427C9A
                                                                                                                                                                   PLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    259 SASIVSPLLE 268
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                                                                                                                                                                                                                                                                                                                                                                                                   532, Application US/09483672A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FastSEQ for Windows Version 3.0
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Retter, Mark
Solk, John
                                                                                                                                                                             Solk, John A.
Day, Craig H.
Skeiky, Yasir A.W.
                                                                                                                                                                                                                                 Kalos, Michael D.
Fanger, Gary R.
Retter, Marc W.
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Mitcham, Jennifer L.
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                                                                                                                                                                                                                                                                                                                                  Mitcham, Jennifer L.
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                                                                                                  210121.42711C11
                                                                                                                                                                                                                                                                                  , Steven G.
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                                                                                                                                                                                                                                                                                                                   Susan Louise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65.5%;
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Pred. No.
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ORGANISM: Homo sapiens

US-09-483-672A-532

Best Local Similarity Matches 7; Conserv

Conservative

65.5%;

Score 36; Pred. No.

DB 18; 1.1e+02;

Length 292

Query Match Best Local S Matches

Similarity

65.5%;

Score 36; DB 19; Length 292; Pred. No. 1.1e+02;

Conservative

2; Mismatches

Indels

0;

Gaps

0;

0; Gaps

:||: ||||| 259 SASIVSPLLE 268

3 NASVTSPLLE 12

09-534-825A-315

Query Match

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NUMBER OF SEQ ID NOS:
SOFTWARE: FESTSEQ for
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
; ORGANISM: Homo sapiens
US-09-536-857-532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity /u.v
                                                                                                       FILE REFERENCE: 210121.4271
CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 2000-C
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SOFTWARE: FastSEQ fo:
SEQ ID NO 315
LENGTH: 292
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TYPE: PRT
ORCANISM: Homo sapiens
-09-534-825A-315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 315, Application US/09534825A GENERAL INFORMATION:
                                                                                                                           PPLICANT: Day, Craig H.
ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
710171 42712C12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/534,825A
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                                                                                                                                                                                                                                                                                                                                                                 INFORMATION
                                                                       FastSEQ for Windows Version 3.0
                                                                                                                                                                                                    Stolk, John A.
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Reed, Steven G.
Kalos, Michael D.
                                                                                                                                                                                                                  Fanger, Gary R.
Retter, Marc W.
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                                                                                                                                                                                                                                                                                                                                                                           Application US/09536857
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for Windows Version 3.0
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                                                                                                       UMBER: US/09/536,857
2000-03-27
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Pred. No.
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SOFTWARE: FastSEQ for Wi
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
US-09-593-793A-532
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Best Local Similarity
"-+-hes 7; Conserv:
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; Sequence 532, Ap
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APPLICANT: Helper, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
ELLE REFERENCE: 210121.42715C15
CURRENT FILLING DATE: 2000-06-13
CURRENT FILLING DATE: 2000-06-13
CURRENCE SEQ ID NOS: 814
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                                                                                                                                                                                                                                       equence 532,
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Day, Craig H. Vedvick, Thomas SCarter, Darrick Li, Samuel
                                                          Fanger, Gary R. Retter, Marc W. Stolk, John A.
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                                                                                                                                                     Harlocker, Susan L.
                                                                                                                                                                      Mitcham, Jennifer L
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                                                                                                           (alos, Michael D.
                                                                                                                        Jiang, Yuqui
Tenderson, F
                                                                                                                                                                                   Dillon, Davin C.
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Vedvick, Thomas S.
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Mitcham, Jennifer L.
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                                                                                                                        Robert A.
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Pred. No. 1.1e+02;
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PPLICANT: Skeiky, Yasir A. PPLICANT: Hepler, William

Skeiky, Yasir A.W.

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Query Match
Best Local Similarity
Thes 7; Consery:
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CURRENT APPLICATION NUMBER: US/09/651,236
CURRENT FILING DATE: 2000-08-29
NUMBER OF SEQ ID NOS: 865
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo Sapiens
US-09-651-236-532
US-09-657-279-532

US-09-657-279-532

: Sequence 532, Application US/09657279

: GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-09-636-215-532
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Best Local Similarity 70.0
Matches 7; Conservative
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SOFTWARE: FastSEQ for
SEQ ID NO 532
LENGTH: 292
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ITLE OF INVENTION:
LE REFERENCE: 210
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259 SASIVSPLLE 268
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Hepler, William
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Carter, Darrick
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Retter, Marc W.
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ick, Thomas S.
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21,121,427,17(1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Robert A.
                                                                                                                                                                 Score 36; DB 20;
Pred. No. 1.1e+02;
2; Mismatches 1
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Pred. No. 1.1e+02;
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                                                                                                                                                                   Indels
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NUMBER OF SEQ ID NOS: 87
SOFTWARE: FastSEQ for W3
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
US-09-657-279-532
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                                                                                                                      Query Match
Best Local Similarity
7; Conserv
Search completed: June Job time: 514 sec
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CURRENT FILING DATE: 2000-09-06
                                                                                                                                                                                                                                                                                                                                           ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

LE REFERENCE: 210121.427C19
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                                                                   259 SASIVSPLLE 268
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Hepler, William
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Carter, Darrick
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Mitcham, Jennifer
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Kalos, Michael
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                                                                                                                                 Conservative
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            4, 2001, 12:20:01
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70.0%;
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                                                                                                                                               Score 36; DB 20;
Pred. No. 1.1e+02;
                                                                                                                                                              Length 292;
                                                                                                                                Indels .
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect.score:
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                         Score
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000
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                                         US-09-232-773A-12

US-09-232-773A-6

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US-09-232-773A-14

US-09-426-371-3

US-09-270-849B-181441

US-09-270-849B-187805

US-09-270-849B-182930

US-09-270-849B-182930

US-09-270-849B-182930
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US-09-590-583-315

US-09-825-301-4

US-09-825-301-532

US-09-439-313-532

US-09-439-313-376

US-09-825-301-2

US-09-825-301-2

US-09-835-301-2

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                US-09-270-849B-182930
US-09-270-849B-183902
US-09-270-849B-181440
PCT-US01-01350-117
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Sequence 299, App
Sequence 376, App
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2, Appli
187805,
344, App
182930,
183902,
183902,
181440,
117, App
2260, Ap
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3, Appli
181441,
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8401, Ap
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376, App
8794, Ap
10537, A
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54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	54.5	
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PCT-US01-04098A-2979	PCT-US01-04098A-1011	PCT-US00-35017A-1259	US-09-816-494-2	US-09-833-245-1202	PCT-US01-11988-1202	PCT-US01-04098A-3285	PCT-US01-04098A-1317	US-09-270-849B-182179	PCT-US01-04098A-3640	PCT-US01-04098A-1672	US-09-640-211A-2277	US-09-739-449-10533	US-09-270-849B-181024	US-09-833-245-1199	PCT-US01-11988-1199	US-09-270-849B-183119	US-09-270-849B-192928	
Sequence 2979, Ap	Sequence 1011, Ap	Sequence 1259, Ap	Sequence 2, Appli	Sequence 1202, Ap	Sequence 1202, Ap	Sequence 3285, Ap	Sequence 1317, Ap	Sequence 182179,	Sequence 3640, Ap	Sequence 1672, Ap	Sequence 2277, Ap	Sequence 10533, A	Sequence 181024,	Sequence 1199, Ap	Sequence 1199, Ap	Sequence 183119,	Sequence 192928,	

### ALIGNMENTS

09-284-107-21

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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Logtenberg, Ton
APPLICANT: de Kruif, Cornelis Adridan John
TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES
TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FO
APPLICANT: Reed, Steven G.
APPLICANT: Smith, John M.
APPLICANT: Misher, Linda E.
APPLICANT: Dillon, Davin C.
APPLICANT: Dillon, Davin C.
APPLICANT: Retter, Marc W.
APPLICANT: Wang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Sheiky, Yasir A.W.
APPLICANT: APPLICANT AND METHODS FOR THE TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF BREAST CANCER
                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                 APPLICANT: Frudakis, Tony N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
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TYPE: PRT
ORGANISM: Artificial Sequence
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Pred. No. 0.00025;
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FILE REFERENCE: 210121.419C9

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JS-09-825-301-4
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Best Local Similarity
7; Conserva
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JS-09-822-827-532
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US-09-590-583-315
                                                           CURRENT APPLICATION NUMBER: US/09/825,301
CURRENT FILING DATE: 2001-04-02
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4
TYPE: PRT
ORGANISM: Homo sapiens
-09-825-301-4
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Best Local Similarity
Matches 7; Conser
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
LENGTH: 292
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SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 315
                                                                                                                                                TITLE OF INVENTION: METHOD TITLE OF INVENTION: AND METHOD FILE REFERENCE: 210121.513
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GENERAL INFORMATION:
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CURRENT FILING DATE: 2000-06-08
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                                                  LENGTH: 292
                                                                                                                                                                                                                                                                                                 PPLICANT: Houghton, Raymond
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND ITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                           259 SASIVSPLLE 268
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                                                                                                                                                           T: Zehentner, Barbara
T: Persing, David H.
TIVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION INVENTION: AND MONITORING OF BREAST CANCER
                                                                                                                                                                                                                                                                                                                                  Application US/09825301
                                                                                                                                                                                                                                                             Molesh, David A.
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Pred. No.
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Pred. No. 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 532
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapiens
US-09-439-313-532
US-09-590-583-299
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Best Local Similarity
"hes 7; Conservations
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Best Local Similarity
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                                                              SEQ ID NO 299
                                                                                                                                                                                                                                                                                                                                                                                      -09-590-583-299
                                                                               NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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                                                                                                 APPLICANT: SKEİKY, YASİT A.W.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: THERAPY AND DIAGNOSIS OF BREAST CANCER
FILE REFERENCE: 210121.419C9
CURRENT APPLICATION NUMBER: US/09/590,583
CURRENT APPLICATION ONDHER: 2000-06-08
CURRENT APPLICATION ONDHER: 2000-06-08
NUMBER OF SEQ ID NOS: 324
           ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C9
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                                                ENGTH:
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                                                329
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Dillon, Davin C.
Retter, Marc W.
Wang, Aijun
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                                                                                                                                                                                                                                                                                            Smith, John M.
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Retter, Mark
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                             Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.5%;
70.0%;
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Pred. No.
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Pred. No.
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Matches

3 NASVTSPLLE 12

Local

l Similarity 7; Conserv

Conservative

65.5%; 70.0%;

Score 36; Pred. No.

DB 5;

Length 329;

Mismatches

Indels

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Gaps

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RESULT 9
US-09-439-313-376
; Sequence 376, Application US/09439313
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CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 376, Application US/09822827 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 2 LENGTH: 329
                                                                                                                                                               Best Local Similarity 70.0 Matches 7; Conservative
                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Homo sapien -09-825-301-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/825,301
CURRENT FILING DATE: 2001-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.534C1
                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Persing, David H.

TITLE OF INVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION FITLE OF INVENTION: AND MONITORING OF BREAST CANCER PRICE REFERENCE: 210121.513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 329
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPLICANT:
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296 SASIVSPLLE 305
                                                                                            296 SASIVSPLLE 305
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nes 7; Conservative
                                                                                                                              3 NASVTSPLLE 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Houghton, Raymond L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zehentner, Barbara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jiangchun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  David A.
                                                                                                                                                                              65.5%;
70.0%;
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70.0%;
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7. Mismatches
                                                                                                                                                                               Score 36; Pred. No.
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Pred. No.
                                                                                                                                                                 Mismatches
                                                                                                                                                                                 DB 5;
43;
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US-09-739-449-10537
Sequence 10537, Application US/09739449
GENERAL INFORMATION:
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Best Local Similarity
"heres 7; Conserv
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                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Agrobacterium tumefaciens US-09-739-449-8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 10
US-09-739-449-8794
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Best Local Similarity 5...
Ches 6; Conservative
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SOFTWARE: FastSEQ for Wind
SEQ_ID_NO_376
                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8794, Application US/09739449
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 199-11-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
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Retter, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dillon, Davin C.
Mitcham, Jennifer L.
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Pred. No.
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/739,449

CURRENT FILING DATE:

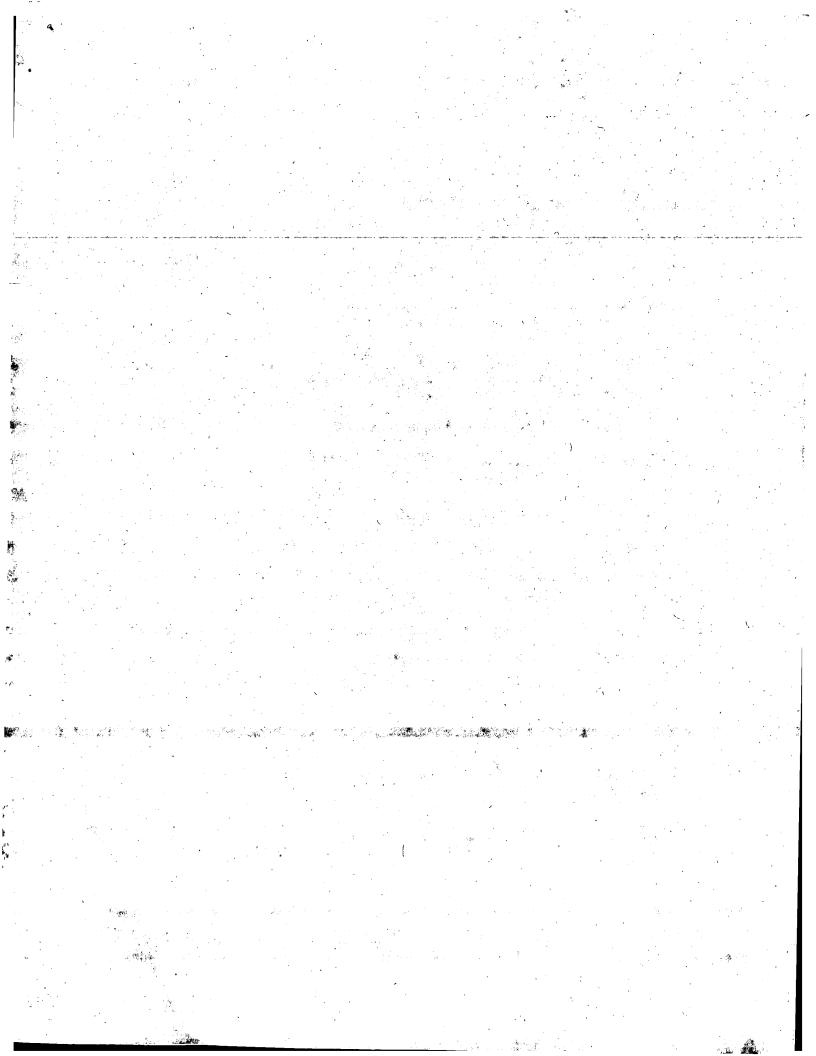
2000-12-19

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FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-270-849B-194755
                                                                                                                                                                                                                                                                                                                                                      RESULT 13
US-09-270-849B-194755
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Query Match
Best Local Similarity bu.v
Thes 6: Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: at any location having Xaa, Xaa means any amino acid US-09-270-849B-189198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-270-849B-18919B; Sequence 18919B; Application US/09270849B; GENERAL INFORMATION:
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                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 194755
                                                                                                                                                                                                                                                                                                        Sequence 194755, Application US/09270849B
GENERAL INFORMATION:
APPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 58.
Best Local Similarity 60.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 189198
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Insect genome survey devices
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Best Local :
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PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID 10537
                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                  FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Agrobacterium tumefaciens -09-739-449-10537
                                                                                                                                                                                       LENGTH: 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                           127
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31 VLNAFYTSPV 40
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                                      58.2%;
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70.0%;
                                    Score 32;
Pred. No.
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Pred. No. 1.2e+02;
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Pred. No. 73;
                       Mismatches
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                                    2.7e+02;
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                                                 Length 341;
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Search completed: June 4, Job time: 590 sec

2001, 12:21:28

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                                       Query Match
Best Local Similarity
---hes 6; Conserv
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US-09-232-773A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/232,773A
CURRENT FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 18
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 12
LENGTH: 661
TYPE: PRT
ORGANISM: Homo sapien
US-09-232-773A-12
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SEQ ID NO 6
LENGTH: 687
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Best Local
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CURRENT FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 12, Application US/09232773A
                                                                                                                                                                                                                                                             TITLE OF INVENTION: INHIBITOR OF VASCULAR ENDOTHELIAL CELL TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                   APPLICANT: Kendall, Richard L. APPLICANT: Thomas, Kenneth A.
                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: 188
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249 VLNCTATTPL 258
                                                                                                                                                                                                                                                                                                                                                                                                                   223
                         1 VLNASVTSPL 10
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                                                    Conservative
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INHIBITOR OF VASCULAR ENDOTHELIAL CELL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GROWTH FACTOR
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                                                                58.2%;
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                                                               Score 32; I
Pred. No. 5
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                                                      Mismatches
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GNLVTLSCETKL 12
                                                                                                                                                                                   Length DB
                                                                                                                                                                                                                                                                                                                                            /cgn2_6/ptodata/2/paa/US60_COMB.pep:
                                                                                                                            US-60-160-209-3193
US-60-160-209-4252
                                                                                                           US-60-160-203-4414
                                                                                                                                                                                                                                      SUMMARIES
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7.418 Million cell updates/sec
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Sequence 3193, Ap
Sequence 4414, Ap
Sequence 7, Appli
Sequence 114, Appl
Sequence 118, Appl
Sequence 10, Appl
Sequence 3220, Ap
Sequence 3220, Ap
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39	39	39	39	39	39	39	39.	39	39	40	42	42	42	42	43	43	43	43	43	46	47	47	48	48	48	48	48	48	48	48	48	48	48	5
63.9	ω.	Ψ	Ψ	Ψ	ω.		ω.	63.9	63.9	5	68.9	68.9	68.9	68.9	70.5	70.5	70.5	70.5		75.4	77.0			78.7						78.7		78.7		1 0 1
147	147	147	147	95	95	95	95	95	59	123	359	359	359	358	359	217	181	181	83.	59	96	59	410	370	370	370	370	370	343	305	278	177	177	2
щ.	_	_	ب	23	23	23	23	21	23	23	19	_	<u> </u>	16	17	_	ᆫ	<u>,</u>	23	23	23	23	ω	21	21	21	21	21	21	21	21	23	23	D
PCT-US01-01349-785	4	-US01-01326-	PCT-US01-01326-158	7-873-	187-470-	169	7-499-53	9-731-872-	-307-	73-469-	US-09-528-003-116	PCT-US00-29363-134	PCT-US00-07285-116	US-09-206-647-1			01349-	PCT-US01-01326-137	US-60-195-053	SD	US-60-194-508-	US-60-170-374-	US-07-861-895-25	US-09-702-021-3	US-09-702-021-	US-09-702-021-3	. US-09-702-021-	US-09-702	US-09-702-021-	US-09-702-021-	-Sn	US-60-185-3	US-60-185-361-731 "	116-09-463-073-
Sequence 785, App	587,	217,	Sequence 158, App	18502	676,	676, 1	5383,	364, 1	2040,	1595,	e 116,	,	116,	Sequence 1, Appli	e 45	664,		137, Ag	1821,	2665,		e 2541,	25, A	Sequence 33, Appl	32,	31,	30,	ω	5, 7	Sequence 34, Appl	6, Ac	e 692,	Sequence 731, App	0 / 777

## ALIGNMENTS

CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3193
LENGTH: 58
TYPE: PRT
ORGANISM: HUMAN
US-60-160-209-3193 US-60-160-209-4252 : Sequence 4252, Application US/60160209 ; GENERAL INFORMATION: 밁 Ş US-60-160-209-3193; Ap Query Match Best Local ( GENERAL INFORMATION:
APPLICANT: BONAZZI, VIVIEN Matches FILE REFERENCE: CL00011 TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES TITLE OF INVENTION: THEREOF Local Similarity nes 12; Conserv 1 GNLVTLSCETKL 12 GNLVTLSCETKL 12 100.0%; Score 61; DB 23; ilarity 100.0%; Pred. No. 0.0021; Conservative 0; Mismatches 0 Application US/60160209 DB 23; Length 58; Indels 0 Gaps

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US-60-160-203-4414
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Matches 12; Conser
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SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4252
LENGTH: 96
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TITLE OF INVENTION: P
TITLE OF INVENTION: P
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SOFTWARE: FastSEQ for
SEQ ID NO 4414
LENGTH: 102
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Best Local :
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Best Local Similarity 100.0%;
Matches 12; Conservative 0
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CURRENT FILING DATE: 1999-10-19
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TITLE OF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: VARIANT
LOCATION: (1)...(96)
OTHER INFORMATION: Xaa = Any Amino Acid
                                              APPLICANT: Epa, Vidana
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF
TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: HUMAN
CURRENT APPLICATION NUMBER: US/09/245,764
CURRENT FILING DATE: 1999-02-05
                                                                                                    APPLICANT: Maxwell, Kelly F. APPLICANT: Garrett, Thomas P.J.
                                                                                                                                       APPLICANT: Hogarth, P. Mark
APPLICANT: Powell, Maree S.
APPLICANT: McKenzie, Ian F.C.
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                                   FILE REFERENCE:
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Pred. No. 0.0036;
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Pred. No.
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US-09-702-021-14
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LENGTH: 261
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Best Local
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SEQ ID NO 14
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                                                                                                                              NUMBER OF SEQ ID NOS: 1694
SOFTWARE: PatentIn Ver. 2.
SEQ ID NO 1128
LENGTH: 399
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Best Local (
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CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
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                                                                                                                                                                                             CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 37
                                                                                                                                                                                                                                                                                                 APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
                             NAME/KEY: SITE
LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                ORGANISM: Homo sapiens
                                                                                FEATURE:
AME/KEY: SITE
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nes 12; Conserv
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Pred. No.
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Pred. No. 0.014;
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; LOCATION: (349); OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids PCT-US00-05882-1128

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RESULT 9
US-60-170-374-2865
; Sequence 2865, Application US/60170374
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US-09-702-021-10
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Best Local Similarity 75.0
Conservative
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Best Local S
Matches 12
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                                                                                                                                              Matches
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                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3220
LENGTH: 75
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                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3220,
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TYPE: PRT
ORGANISM: Homo sapiens
-09-702-021-10
                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/60/170,373
CURRENT FILING DATE: 1999-12-13
NUMBER OF SEQ ID NOS: 4282
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CURRENT FILING DATE: 2000-10-30
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ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                          ILE REFERENCE:
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Local Similarity 100.0%;
hes 12; Conservative (
                                                                                                                                            Local Similarity 75.
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75.08;
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75.08;
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Pred. No. 0.42
2; Mismatches
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Pred. No.
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Pred. No.
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Best Local Similarity
Matches 9; Conser
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CURRENT FILING DATE: 1999-12-13
NUMBER OF SEQ ID NOS: 3666
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ISOLATE TITLE OF INVENTION: ACID MOI TITLE OF INVENTION: THEREOF FILE REFERENCE: CLOOO146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                 INFORMATION FOR SEQ ID NO:
              SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Relicurrent APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                   REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
                                                                                                                                                                                                                                        CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 07/896,457
FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: FILING DATE: 31-00
                                                                                                                                                                                        NAME: BENT, Stephen A REGISTRATION NUMBER: :
                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington
                                                                                                                   TELEFAX:
                                                                                                                                 TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 GSLVTLRCQTKL 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      20007-5109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/08332562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3: Foley & Lardner
3000 K Street, N.W.,
                                                                                                                     (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HULETT, Mark D.
VENTION: FC.RECEPTOR FOR IMMUNOGLOBULIN EQUENCES: 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HIBBS, Margaret L.
                                                                                                                                                                                                                                                                                                                                                               PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HOGARTH, Mark P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 McKENZIE, Ian F.C.
linear
                                                                                                                                                                                                                                                                                                         31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ISOLATED HUMAN KINASE PROTEINS, NUCLEIC AND USES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bernadette M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78.78;
75.08;
                                                                                                                                                                                                                                                                                                                             US/08/332,562
                                                                                                                                                                                    29,768
                                                                                   94:
                                                                                                                                                                      54270/119/GRHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 48;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Suite 500
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0.54;
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Query Match Best Local

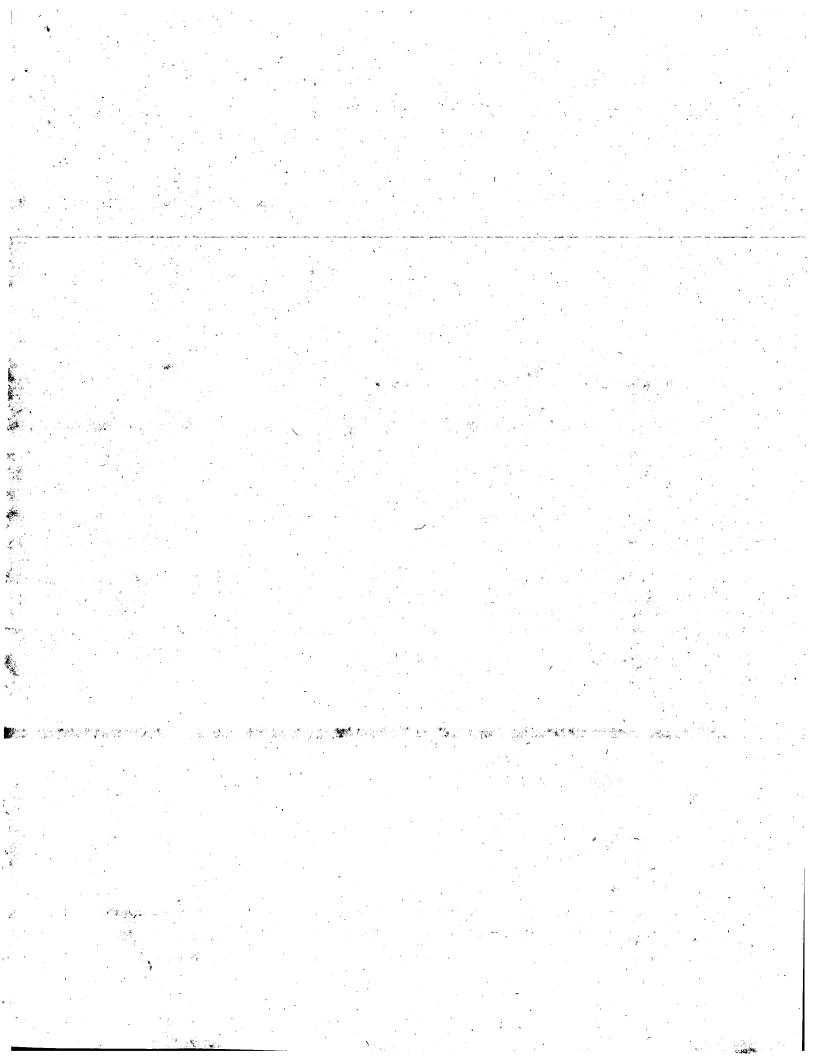
78.7%; 75.0%;

Length 91;

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Query Match
Best Local Similarity
---hes 9; Conserv:
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                                                                                                                                                                            NFORMATION FOR SEQ ID NO:
                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
                                                                                                                                                                               TELLEFAX: 1-16
TELEFAX: 904136
                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 16
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: AU PH6166
FILING DATE: 26-MAY-1986
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 1
FILING DATE: 29-MAY-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US 08/332/562
FILING DATE: 31-OCT-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
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                                                                                                                  STRANDEDNESS:
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                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NPPLICANT: HULETT, MARK D.
TITLE OF INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
TUMBER OF SEQUENCES: 130
                                                                                                                                                                                                                                                             NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 27-MAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 05-JU
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 GSLYTLNCETNL 28
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3000 K Street, N.W.,
                                                                                                   linear
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HIBBS, Margaret L.
SCOTT, Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BONADONNA, Lisa
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                                                                                                                                                                                                                                                                                                                                                                                                  27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                            16-SEP-1994
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                                                                                                                 single
                          78.7%;
75.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/462,973
                                                                                                                                                                            . 94 :
   Score 48; DB 8; Le
Pred. No. 0.66;
2; Mismatches '1;
                                                                                                                                                                                                                                                  16786/203
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Pred. No. 0.66;
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                                   DB 8; Length 91;
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         Indels
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US-09-702-021-6
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Best Local Similarity
'--hes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                     ; ORGANISM: HUMAN US-60-185-362-692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  В
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US-60-185-361-731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-60-185-362-692
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                                                                                                   Sequence 6, Application US/09702021 GENERAL INFORMATION:
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APPLICANT: C. Fraser
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF
FILE REFERENCE: 7853-217.
CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 836
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 692, Applic GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 731 LENGTH: 177 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION: APPLICANT: Bonazzi,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/60/185,362
CURRENT FILING DATE: 2000-02-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 731,
                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 177
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: ISOLATED HUMAN DRUG TARGET PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN DRUG TARGET PROTEINS.
TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: CL000285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bonazzi, Viven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/60/185,361
CURRENT FILING DATE: 2000-02-28
NUMBER OF SEQ ID NOS: 968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS, AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000291
                                                                                                                                                                                                                                     1 GNLVTLSCETKL 12
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                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                         78.7%;
75.0%;
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                                                                                                                                                                                                                                                                                                         Score 48; |
Pred. No. 1
                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                       DB 23; Length 177;
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; SOFTWARE: FastSEQ for Wi
; SEQ ID NO 34
; LENGTH: 305
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-702-021-34
Search completed: June 4, 2001, 12:20:01 Job time: 514 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 15
US-09-702-021-34
US-09-702-021-34
; Sequence 34, Application US/09702021
; SEQUENCE 34, Application US/09702021
; GENERAL INFORMATION:
APPLICANT: C. Fraser
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF
FILE REFERENCE: 7853-217
; CURRENT APPLICATION NUMBER: US/09/702,021
; CURRENT FILING DATE: 2000-10-30
; NUMBER OF SEQ ID NOS: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 45
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 6
LENGTH: 278
TYPE: PRT
ORGANIZM: Homo sapiens
US-09-702-021-6
                                                                                                                                                                                                     Query Match 78.7%; Score 48; DB 21; Length 305; Best Local Similarity 75.0%; Pred. No. 2.2; Matches 9; Conservative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 78.7%;
Best Local Similarity 75.0%;
Matches 9; Conservative
                                                                                                             1 GNLVTLSCETKL 12
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120 GSLVTLRCQTKL 131
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93 GSLVTLRCQTKL 104
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Database
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Maximum DB
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
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DB
                                            US-09-514-009-33

US-09-184-467-20

US-09-184-467-24

US-09-184-467-16

US-09-184-467-28

US-09-184-467-28

US-60-248-505-847

US-60-248-505-1220

US-09-270-849B-1220

US-09-270-849B-1563

PCT-US01-11988-1679

PCT-US01-11988-1680
                                                                                                                                                                                                                                          US-09-565-522B-12
US-09-723-716A-43
US-09-723-716A-68
US-09-723-716A-70
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US-09-51
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US-09-270-849B-180519
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AT TONMENTS	US-09-828-825-6	US-09-828-825-16	US-09-828-825-8 .	US-09-828-825-2	US-09-828-825-4	PCT-US00-35017A-1264	PCT-US00-35017A-1153	US-09-832-312-40	US-09-832-312-38	US-09-832-312-36	US-09-832-312-34	US-09-832-312-3	US-09-832-312-5	PCT-US01-10462-13	US-09-832-312-9	US-09-833-245-208	PCT-US01-11988-208	US-08-849-406-38	
	Sequence 6, Appli	Sequence 16, Appl	Sequence 8, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 1264, Ap	Sequence 1153, Ap	Sequence 40, Appl.	Sequence 38, Appl	Sequence 36, Appl	Sequence 34, Appl	Sequence 3, Appli	Sequence 5, Appli	Sequence 13, Appl	Sequence 9, Appli	Sequence 208, App	Sequence 208; App	Sequence 38, Appl	-

#### Query Match Best Local Similarity "~~~hes 12; Conserv ; OTHER INFORMATION: Oligopeptides representing the extracellular; OTHER INFORMATION: domain of CD64 US-09-284-107-22 US-09-284-107-22 NUMBER OF SEQ ID NOS: 29 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 22 Sequence 22, Applica GENERAL INFORMATION: CURRENT APPLICATION NUMBER: US/09/284,107 CURRENT FILING DATE: 1999-10-25 PRIOR APPLICATION NUMBER: PCT/NL97/00557 PRIOR FILING DATE: 1997-10-07 PRIOR APPLICATION NUMBER: EP 96202791.8 PRIOR FILING DATE: 1996-10-08 TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AIFILE REFERENCE: 313632000600 APPLICANT: Logtenberg, Ton APPLICANT: de Kruif, Corn LENGTH: 12 TYPE: PRT ORGANISM: Artificial Sequence FEATURE: Application US/09284107 Cornelis Adriaan John 100.0%; Score 61; Pred. No. DB 5; Length 12; AFFINITY F FOR A TARGET

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1 GNLVTLSCETKL 12 |||||||||| 1 GNLVTLSCETKL 12

Conservative

Mismatches

Indels

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Gaps

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PCT-US01-11988-133 Sequence 133, Application:

Application PC/TUS0111988

APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REFERENCE: PF546PCT

CURRENT APPLICATION NUMBER: PCT/USO1/11988
CURRENT EILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12

PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2000-1

60/256, 931 60/199, 384

; TYPE: PRT; ; ORGANISM: Homo sapiens PCT-US01-11988-133

ENGTH:

ID NO 133

PRIOR FILING DATE: 2000-04-25 NUMBER OF SEQ ID NOS: 2267 SOFTWARE: Patentin Ver. 2.1

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NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 180519
LENGTH: 257
TYPE: PRT
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Best Local Similarity
Anhas 8; Conserv
                                                                                                                                                                                                                                                                              RESULT 4
US-09-270-849B-180519
OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-270-849B-180519
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                                                                                                                                                                                                                                               Sequence 180519, App
GENERAL INFORMATION:
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Best Local
                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/270,849B CURRENT FILING DATE: 1999-03-17
                                                                                                                                                                                                      APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                              FILE REFERENCE:
                                   FEATURE:
                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
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PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RIOR FILING DATE: 2000-12-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     equence 133,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT: Human Genome Sciences, Inc.
ITLE OF INVENTION: Albumin Fusion Proteins
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                                                                                                                                                                                                                                                                                                                                                     185 GGPMTLSCQTKL 196
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66.7%;
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Pred. No.
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Query Match

Score

38;

DB 5;

Length 257;

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; LENGTH: 544
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-565-522B-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-723-716A-43
SEQ ID NO 43
LENGTH: 28
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Best Local Similarity
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                                                                                PRIOR APPLICATION NUMBER: GB 9810759.2 PRIOR TILING DATE: 1998-05-19 PRIOR APPLICATION NUMBER: GB 9821129.5 PRIOR FILING DATE: 1998-09-29
                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/723,716A CURRENT FILING DATE: 2000-11-28 PRIOR APPLICATION NUMBER: US 09/335,087 PRIOR FILING DATE: 1999-06-17 PRIOR APPLICATION NUMBER: PCT/GB99/01588 PRIOR FILING DATE: 1999-05-19 PRIOR FILING DATE: 1999-05-19
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                                          SOFTWARE: PatentIn version 2.1
                                                                                                                                                                                                                                                                                             APPLICANT: WILLCOX, Benjamin Ernest
APPLICANT: BOULTER, Jonathan Michael
TITLE OF INVENTION: Soluble T Cell Receptor
FILE REFERENCE: 102286.499DIV
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PRIOR FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 12
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PRIOR FILING DATE: 2000-04-25
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APPLICANT: Mize, Nancy K
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TILE OF INVENTION: Methods and Materials Relating to Novel
TILE OF INVENTION: Polypeptides and Polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LE REFERENCE: HYS-8
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                                                                                                                                                                                                                                                                                                                                                                                GAO, George Fu
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Liu, Chenghua
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1; Mismatches
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                                                                                                                                                                                       JS-09-723-716A-70
                                                                                                                                           Sequence 70, Application US/09723716A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 68
LENGTH: 287
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Best Local Similarity
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SOFTWARE: PatentIn ver
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                                                                                                        APPLICANT:
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PRIOR APPLICATION NUMBER: PCT
PRIOR FILING DATE: 1999-05-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BOULTER, Jonathan Michael ITILE OF INVENTION: Soluble T Cell Receptor TILE REFERENCE: 102286.409DIV
"URRENT APPLICATION NUMBER: US/09/723,716A
"URRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION:
                                                                                                                        APPLICANT: JAKOBSEN, Bent Karsten
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Amino acid OTHER INFORMATION: sequence of human HLA-A2/flu matrix peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RIOR APPLICATION NUMBER: US 09/335,087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: Predicted amino acid sequence of the soluble OTHER INFORMATION: HLA-AZ/flu matrix restricted beta chain from JM22, OTHER INFORMATION: as fused to the "lecine zipper" domain of c-fos. OTHER INFORMATION: (Figure 11).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                        ITLE OF
                                                                                                                                                                                                                                                                          18 GONVTLSCEONL 29
                                                                                                                                                                                                                                                                                                               1 GNLVTLSCETKL 12
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T: WILLCOX, Benjamin Ernest
T: BOUTTER, Jonathan Michael
INVENTION: Soluble T Cell Receptor
ERENCE: 102286.409DIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PatentIn version 2.1
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                                                                              BELL, John Irving
GAO, George Fu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 leucine zipper domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence of human HLA-A2/flu matrix peptide restricted JM22 TCR beta chain fused to c-fos
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                Length 287;
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                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Mus musculus US-09-514-009-46
                                                                                                                                               US-09-514-009-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-514-009-46
Sequence 36, Application US/09514009
GENERAL INFORMATION:
APPLICANT: Fraser, Christopher C.
TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF
FILE REFERENCE: 07334-251001
CURRENT APPLICATION NUMBER: US/09/514,009
CURRENT FILING DATE: 2000-02-25
                                                                                                                                                                                                                                                                                                             Query Match.

Best Local Similarity

Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 46, Application US/09514009 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 8; Conserval
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF FILE REFERENCE: 07334-251001
CURRENT APPLICATION NUMBER: US/09/514,009
CURRENT FILING DATE: 2000-02-25
PRIOR APPLICATION NUMBER: US 09/259,387
PRIOR FILING DATE: 1999-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: GB 9821129.5
PRIOR FILING DATE: 1998-09-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR EILING DATE: 1998-05-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/723,716A
CURRENT FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Fraser, Christopher C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 1999-05-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT/GB99/01588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-06-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Amino acid OTHER INFORMATION: sequence of human HLA-A2/flu matrix peptide OTHER INFORMATION: restricted JM22 TCR beta chain fused to c-fos OTHER INFORMATION: leucine zipper domain and BirA biotinylation tag.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 09/335,087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18 GONVTLSCEONL 29
                                                                                                                                                                                                                                                    1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GNLVTLSCETKL 12
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                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                   57.48;
63.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       60.78;
                                                                                                                                                                                                                                                                                                                  Score 35; DB Pred. No. 20; 1; Mismatches

    Mismatches

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Pred. No.
                                                                                                                                                                                                                                                                                                                                                        Length 57;
                                                                                                                                                                                                                                                                                                             Indels
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PRIOR APPLICATION NUMBER: US 09/259,387 PRIOR FILING DATE: 1999-02-26

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; NAME/KEY: SIGNAL
; LOCATION: (1)...(34)
US-09-514-009-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity b...
hes 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: PRT
; ORGANISM: Mus musculus
US-09-514-009-34
                                                                                                                                                                                                                                                          Sequence 3, Application US/09514009
GENERAL INFORMATION:
APPLICANT: Fraser, Christopher C.
TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF
FILE REFERENCE: 07334-251001
CURRENT APPLICATION NUMBER: US/09/514,009
CURRENT APPLICATION NUMBER: US/09/514,009
                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 63.b
""" 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-514-009-34
                                                                                                                                                                                NUMBER OF SEQ ID NOS: 51

SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 33

LENGTH: 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Mus musculus US-09-514-009-36
                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/259,387 PRIOR FILING DATE: 1999-02-26
                                                                                                                                      TYPE: PRT
ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: SOFTWARE: FastSEQ for SEQ ID NO 36
                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/259,387
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/514,009
CURRENT FILING DATE: 2000-02-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Fraser, Christopher C.
TITLE OF INVENTION: SECRETED PROTEINS AND USES THEREOF
FILE REFERENCE: 07334-251001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                25 GONVSLSCSTK 35
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         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       57.4%;
63.6%;
                       63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         57.4%;
63.6%;
                  Score 35; DB 5;
Pred. No. 77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 35; DB Pred. No. 37;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 5; Length 205;
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                                  Length 239;
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    Indels
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Gaps
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RESULT 14
US-09-184-467-24
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Best Local Similarity
-+~hes 6; Conserv:
                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 20:
US-09-184-467-20
                                                                                                                              Sequence 24, Application US/09184467 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 13
US-09-184-467-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (212) 664-0525
TELEX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acids
                                                                                                                                                                                                                                                  380 GNTITLPCRIK 390
                                                                                                                                                                                                                                                                                      1 GNLVTLSCETK 11
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              59 GONVSLSCSTK 69
                                                                      TITLE OF INVENTION: HIV-1 VACCINES,
                                                                                                              APPLICANT: Hasel, Karl W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/530,146
FILING DATE: <UNKNOWN>
ATTORNEY_RAGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE_DOCKET NUMBER: 41190/JPW/AJM
TELEPHONE: (212) 977-950
TELEPHONE: (212) 977-950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Maddon, Paul J.
TITLE OF INVENTION: HIV-I VACCINES,
THERETO, AND THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hasel, Karl W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/184,467 FILING DATE: 02-Nov-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Cooper & Dunham
STREET: 30 Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid
                                                                                                                                                                                                                                                                                                                                      57.4%;
54.5%;
                                      THERETO,
                                                                                                                                                                                                                                                                                                                                          Pred. No. 1
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                                                      AND THERAPEUTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ES, ANTIBODY COMPOSITIONS RELATED THERAPEUTIC AND PROPHYLACTIC USES
                                                ANTIBODY COMPOSITIONS RELATED ERAPEUTIC AND PROPHYLACTIC USES
                                                                                                                                                                                                                                                                                                                  4; Indels
                                                                                                                                                                                                                                                                                                                                                      Length 479;
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ADDRESSEE: Cooper & Dunham STREET: 30 Rockefeller Plaza

CITY: New York STATE: New York COUNTRY: USA

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/184,467
FILING DATE: 02-Nov-1998
CLASSIFICATION: <dnchown>
PRIOR APPLICATION DATA:

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RESULT 15
US-09-184-467-16
US-09-184-467-16; Sequence 16, Application US/09184467; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 24:
US-09-184-467-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/530,146

FILING DATE: CURK HOWEN
ATTORNEY/AGENT INFORMATION:

NAME: White, John P

REGISTRATION NUMBER: 28,678

REFERENCE/DOCKET NUMBER: 41190/JPW/AJM

TELECOMMUNICATION INFORMATION:

TELEPHONE: (212) 977-9550

TELEPAX: (212) 664-0525

TELEPAX: 422523 COOPUI

INFORMATION FOR SEQ ID NO: 24:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 57.4%;
Best Local Similarity 54.5%;
Matches 6; Conservative
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.24
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/09/184,467
FILING DATE: 02-NOV-1998
CLASSIFICATION LORAND
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/530,146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    380 GNTITLPCRIK 390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HIV-1 VACCINES, ANTIBODY COMPOSITIONS RELATED.
THERETO, AND THERAPEUTIC AND PROPHYLACTIC USES
                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hasel, Karl W
                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Cooper & Dunham
STREET: 30 Rockefeller Plaza.
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 479 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                                                           STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                  THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 35; DB 5;
Pred. No. 1.5e+02;
1; Mismatches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 479;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-184-467-16
                                                                                 망
Search completed: June 4, 2001, 12:21:28
Job time: 590 sec
                                                                                                                                               Query Match 57.

Best Local Similarity 54.

Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                TELEFAX: (212) 664-0525
TELEX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
                                                                                 408 GNTITLPCRIK 418
                                                                                                  1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 977-9550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                57.48;
                                                                                                                                                                Score 35;
Pred. No.

    Mismatches

                                                                                                                                                                                                                                                     16:
                                                                                                                                                                   1.6e+02;
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                                                                                                                                                                                 Length 507;
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Result
No.
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Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Run on:
                                                                                                                                                                                                                                                   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  seq length: 0 seq length: 2000000000
395555561
                                                                                                          100.0
                                                                                                                                                                           Match
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Gapop 10.0 , Gapext 0.5
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61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                June 4, 2001, 12:20:01; Search time 260.2 Seconds (without alignments)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1009251 seqs, 160854530 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                            Length
                                                                                                                                                                            BB
                       3 US-60-160-203-4414

1 US-99-702-021-14

PCT-US00-05882-1128

US-08-332-562-94

US-08-462-973-94

US-08-462-973-94

US-08-61-895-25
                                                                                                                         US-60-160-209-3193
US-60-160-209-4252
                                                                                                                                                                                                                          SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.418 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                   COMB.pep
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                                                                        Sequence 3193, Ap
Sequence 4252, Ap
Sequence 4414, App
Sequence 1128, Ap
                      Sequence 94, Appl
Sequence 94, Appl
Sequence 25, Appl
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	45	44	43	42	41	40	39	38	37	36	<u>3</u> 5	34	ω W	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
-	36	36	37.	37	37	37	37	37	37	37	37	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	39	39	39	39	39	39
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	23	23	16	23	23	23	23	16	16	17	18	16	14	21	17	16	16	14	14	13	12	11	21	17	13	12	11	23	16	23	23	23	23	23	18
	-60-191-681-2158	-191-637-2	-09-252-991A-287	-60-191-681-1814	-60-191-637-22	-60-173-464-1887	-60-161-932-24	-09-252-691C-7	-09-252-691-76	-09-328-352-66	-09-489-039A-9	-09-293-505-1	0-939A-	-09-711-11	98-239-	-09-293-505-1	-09-293-505-	-09-060-939A-1	-09-060-939	3-823-	-80-	-08-720-484-	-09-711-111-	-09-398-239-	8-953-82	-08-897-798-	-08-720-484-	-60-229-515-7	S-09-252-991A-	S-60-191-681-3	S-60-191-637-47	S-60-173-464-40	S-60-167-217-	0-146-055-774	-09-489-0
•	2158	Sequence 26977, A	2877	18143,	22997,	18875,	equence 249	e 7682,	equence 7682,	Sequence 6649, Ap	e 9855,	15	15	7,	e 2,	e :17	e 16,	Ō	16	e 2,	Sequence 7, Appli	e 2,	е 6,	е4,	O	e 6,	e 4, Appl	e 758	e 17591	e 387,	e 477, Ap	e 405, Ap	452, Ap	e 774, Ap	Sequence 13963, A

# ALIGNMENTS

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                                                                                                                                   Qy
                                                                                                                                                                                                                                                          ; ORGANISM: HUMAN US-60-160-209-3193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1
US-60-160-209-3193
US-60-160-209-4252; Sequence 4252, Application US/60160209; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3193
LENGTH: 58
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: BONAZZI, VIVIEN
APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL000113
                                                                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 3193,
                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 1999-10-19
                                                                                                    13
                                                                                                                                      1 LLQRPGLQLYFS 12
                                                                                                  LLQRPGLQLYFS 24
                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/60160209
                                                                                                                                                                                     100.0%;
                                                                                                                                                                                       Score 61; DB 23
Pred. No. 0.001;
                                                                                                                                                                         Mismatches
                                                                                                                                                                                                  DB 23; Length 58;
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US-09-702-021-14

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Вb
                              NUMBER: Fas
SOFTWARE: Fas
SEQ ID NO 14
"FNGTH: 374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: VARIANT
LOCATION: (1)...(96)
OTHER INFORMATION: Xaa = Any Amino Acid
US-60-160-209-4252
                                                                                                                                                                                      Sequence 14, Application US/09702021 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 4414
LENGTH: 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4414, Application US/60160203 GENERAL INFORMATION:
                                                                                                   CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
                                                                                                                   APPLICANT: C. Fraser
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF
FILE REFERENCE: 7853-217
                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: HUMAN
-60-160-203-4414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                   NUMBER OF SEQ ID NOS:
                     LENGTH: 37
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/160,203 CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 4252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 96
TYPE: PRT
ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 4646
SOFTWARE: FastSEQ for Windows Version 4.0
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: ISOLATE:
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: CL000113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BONAZZI,
                                                                                                                                                                                                                                                                                      31 LLQRPGLQLYFS 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INVENTION: NUCLEIC ACID MOLECULES INVENTION: USES THEREOF
                                                               FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ISOLATED HUMAN SECRETED PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISOLATED HUMAN KINASE PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                      Score 61; DB 23;
Pred. No. 0.0018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.0017
                                                                                                                                                                                                                                                                                                                                                                                          DB 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENCODING HUMAN SECRETED PROTEINS AND
                                                                                                                                                                                                                                                                                                                                                                                    Length 102;
                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 96;
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RESULT 6
US-08-332-562-94
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Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1128, Application PC/TUS0005882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
                                                                                                                                                                                                                                                                                                            ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -US00-05882-1128
                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: SITE LOCATION: (349) OTHER INFORMATION: Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
NAME/KEY: SITE
                                                                                                STREET: 5000
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENGTH:
                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                         242 LLQRPGLQLYFS 253
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                 COUNTRY:
                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    217 LLQRPGLQLYFS 228
                                                                                                                                                                                                                                                                                                                                                                                                                           1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 LLQRPGLQLYFS 12
                                                                 20007-5109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12; Conservative
                                                                                                                                                                                                INVENTION:
                                                                                                                                                                                                                                                                                                                           Application US/08332562
                                                                                                                              3: Foley & Lardner
                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                        MCKENZIE, Ian F.C.
HOGARTH, Mark P.
HIBBS, Margaret L.
                                                                                                                                                                                                            HULETT, Mark D.
                                                                                                                                                                                                                           BONADONNA, Lisa
                                                                                                                                                                                                                                             Bernadette M.
                                                                                                                                                                             FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          equals any of the naturally occurring L-amino acids
                                                                                                                            N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 61; DB 21;
Pred. No. 0.0069;
                                                                                                                              Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.0074;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 374;
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Query Match
Best Local Similarity
Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quence 94, Application
                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPETATING SYSTEM: PC-DOS/MS-DOS
                                                              PRIOR APPLICATION DATA:
                                                                             RIOR APPLICATION DATA:
APPLICATION NUMBER: AU PM8232/94
FILING DATE: 16-SEP-1994
                                                                                                                                                           RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT:
                                                                                                                                                                                                                         URRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29 LLQRPGLQLHFS 40
                                                                                                                            APPLICATION NUMBER: US 0 FILING DATE: 31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                         DRRESPONDENCE ADDRESS:
                                               APPLICATION NUMBER:
                                                                                                                                                                            CLASSIFICATION: 435
                                                                                                                                                                                           APPLICATION NUMBER: US/0
FILING DATE: 05-JUN-1995
                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                              ZIP: 20007-5109
                                                                                                                                                                                                                                                                                                                                                         CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 LLQRPGLQLYFS 12
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                                                                                                                                                                                                                                                                                                                                                                                         E: Foley & Lardner
3000 K Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91 amino acids
                                                                                                                                                                                                                                                                                                                                            USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   HULETT, MARK D.

JENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HIBBS, Margaret L. SCOTT, Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HOGARTH, Mark P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MCKENZIE, Ian F.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BONADONNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         UMBER: US 07/896,457
27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-OCT-1994
                                 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
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91.7%;
                                                                                                                                             us 08/332,562
                                                                                                                                                                                                                                                                                                                                                                                                                                            130
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                                               US 07/896,457
WO PCT/AU87/00159
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Pred. No. 0.013;
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                                                                                                                                                                                                                                                                                                                                                                                           Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 904136
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
TELEX: 904136
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                              FILING DATE: 25-OCT-PRIOR APPLICATION DATA:
                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                        REFERENCE/DOCKET NUMBER: 17
ELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 LLQRPGLQLYFS 12
                          TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                          APPLICATION NUMBER: U: FILING DATE: 19920624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: AU E
                                          TELEPHONE:
                                                                                                                                   APPLICATION NUMBER: FILING DATE: 25-0C1
                                                                                                                                                                            APPLICATION NUMBER: PCT/1
FILING DATE: 25-OCT-1990.
                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: BENT, Stephen A. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                             Washington, D.C.
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                                                                                                                                                                                                                                                                                                                                                                                                             3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   : (202)672-5300
(202)672-5399
                            (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HULETI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MCKENZIE,
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IERINO, Francesco L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HOGARTH, Phillip M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                            (202)672-5300
                                                                                                                             NUMBER: AU PJ 7045
25-OCT-1989
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91.7%;
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                                                                                                                                                                                             PCT/AU90/00513
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Pred. No.
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0.013;
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TYPE: AMINO ACID

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; MOLECULE TYPE; protein US-07-861-895-25
                                                                                                                  CURRENT APPLICATION NUMBER: US/60/146,394
CURRENT FILING DATE: 1999-07-29
NUMBER OF SEQ ID NOS: 1866
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1032
LENGTH: 373
                                                                                                                                                                                                                                                                             Sequence 1032, Application to GENERAL INFORMATION:
APPLICANT: BONAZZI, Vivien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 91.7
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 7
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                      60-146-394-1032
                                                                                                                                                                                                                            TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: CLOO
FEATURE:
NAME/KEY: VARIANT
LOCATION: (1)...(373)
OTHER INFORMATION: Xaa =
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-245-764-7
                                                                               ORGANISM: Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   URRENT APPLICATION NUMBER: US/09/245,764
URRENT FILING DATE: 1999-02-05
ARLIER APPLICATION NUMBER: 60/099/994
ARLIER FILING DATE: 1998-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ARLIER APPLICATION NUMBER: 60/073,972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT: Hogarth, P. Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                               198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   232 LLQRPGLQLHFS 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LICANT: Maxwell, Kelly F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LICANT: Powell, Maree S. LICANT: McKenzie, Ian F.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                  1 LLQRPGLQLYFS 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/09245764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Garrett, Thomas P.J.
                                                                                                                                                                                                                                                                                                                     Application US/60146394
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                                                                                                                                                                                                                                                                                                                                                                                                             209
                                                                                                                                                                                                                                         ISOLATED DROSOPHILA PROTEINS, NUCLEIC ACID MOLECULES ENCODING DROSOPHILA PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90.2%;
91.7%;
       Any Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 55; 1
Pred. No. 0
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                66;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 261;
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NAME/KEY: VARIANT
LOCATION: (1)...(611)
OTHER INFORMATION: Xaa - Any Amino Acid
US-60-146-055-774
                                                                                                                                                                                                                                                                                                                                                                                                US-60-146-055-774; Sequence 774; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Klebsiella pneumoniae US-09-489-039A-13963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                 Matches
                                                 Query Match
Best Local :
                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 938
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 774
LENGTH: 611
                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 63.9
Best Local Similarity 80.0
Matches 8; Conservative
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SEQ ID NO 13963
LENGTH: 535
TYPE: PRT
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                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/60/146,055 CURRENT FILING DATE: 1999-07-28
                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: ISOLATED SECRETED PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -09-489-039A-13963
                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: CL000067
                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bonazzi, Vivien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
                                                                                                                                                                                       TYPE: PRT
ORGANISM: Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Gary Breton et. al
                                                                                                                                                                            FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            116 LLQRYGMQLY 125
                                                Local
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3 QRPGLQLYF 11
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                                  Similarity 7; Conser
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                               Conservative
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PNEUMONIAE FOR DIAGNOSTICS A
                                              63.9%; Score 39; 77.8%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        63.9%;
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7.7.8%;
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'Pred. No. 77;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 39; DB 18;
Pred. No. 1.1e+02
                               Mismatches
                                              DB 23;
1.3e+02;
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77;
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                                                                                                                                                                                                                                                                                                                               PROTEINS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                              NUCLEIC ACID
                                                           Length 611;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCES RELATING TO KLEBSIELLA
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                             Indels
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                             Gaps
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414 ERPDLQLYF 422

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US-60-173-464-405

Sequence 405, Application US/60173464

GENERAL INFORMATION:

APPLICANT: Li, Peter W.D.

TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING GPCR PROTEINS AND USES

TITLE OF INVENTION: HERBOF

FILE REFERENCE: CL000173

CURRENT APPLICATION NUMBER: US/60/173,464

CURRENT FILING DATE: 1999-12-29

NUMBER OF SEQ ID NOS: 30269

SOFTWARE: FASTSEQ for Windows Version 4.0

SEQ ID NO 405

LENGTH: 612

TYPE: PRT
ORGANISM: Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   В
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US-60-167-217-452
                                                                                                                                                                                                                                                                                   RESULT 15
US-60-191-637-477
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 63.9
Best Local Similarity 77.1
Matches 7; Conservative
                                  Sequence 477, Application US/60191637
GENERAL INFORMATION:
APPLICANT: Venter, J. Craig
TITLE OF INVENTION: NUCLEIC ACID DETECTION KITS COMPRISING
TITLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM THE DROSOPHILA GENOME, AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000392
CURRENT APPLICATION NUMBER: US/60/191,637
CURRENT APPLICATION NUMBER: US/60/191,637
CURRENT FILING DATE: 2000-03-23
NUMBER OF SEQ ID NOS: 42660
SOFTMARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 63.9
Best Local Similarity 77.8
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/60/167,217
CURRENT FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 23195
SOFTWARE: FastSEQ for Windows Version 4.0
1EQ ID NO 452
LENGTH: 612
TYPE: PRT
EQ ID NO 477
LENGTH: 612
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RPPLICANT: L1, Peter W. D.
TITLE OF INVENTION: ISOLATED DROSOPHILA PROTEINS, NUCLEIC
TILE OF INVENTION: ACID MOLEUCLES ENCODING DROSOPHILA PROTEINS, AND USES
TITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                     3 QRPGLQLYF 11
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415 ERPDLQLYF 423
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77.88;
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77.8%;
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Pred. No.
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Pred. No. 1
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Search completed: June 4, Job time: 515 sec

2001, 12:20:02

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Result
No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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seq.length: 2000000000
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61
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     522552255
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Copyright (c) 1993 - 2000 Comp
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/cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
US-09-828-498-2
US-09-739-449-11020
US-09-74-475-3
US-09-769-066-21
US-09-769-08-98-11176
US-09-270-84.98-189872
US-09-270-84.98-12557,
PCT-US01-11988-1481
PCT-US01-11988-1482
US-09-685-330-6
US-09-685-330-6
US-09-685-330-6
US-09-685-330-6
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US-09-833-245-1481
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US-09-833-245-1481
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US-09-311-62
US-09-270-84
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US-09-7
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US-09-270-849B-188156
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PCT-US01-11988÷1864
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12557, A
1481, Ap
1482, Ap
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11176, A
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11020, A
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186298,
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,	Sequence 193923,	Sequence 8238, Ap	Sequence 180893,	Sequence 1125, Ap	Sequence 182901,	Sequence 195, App	•	Sequence 988, App	•	Sequence 163, App	Sequence 22, Appl	Sequence 6, Appli	Sequence 150, App	Sequence 149, App	Sequence 6, Appli	Sequence 180, App	Sequence 38, Appl	Sequence 1125, Ap

### ALIGNMENTS

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OTHER INFORMATION: Oligopeptides representing the extracellular OTHER INFORMATION: domain of CD64 US-09-284-107-23
                                                                                                                                                                                                                              US-09-270-849B-188156; App
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Best Local Similarity lvv.
12; Conservative
                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect
CURRENT APPLICATION NUMBER: US/09/270;849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 188156
LENCTH 259
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Logtenberg, Ton
APPLICANT: de Kruif, Cornelis Adriaan John
TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES
TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FOR A TARGET
FILE REFERENCE: 313632000600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
                                                                                                                                               FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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1 LLQRPGLQLYFS 12
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                                                                                                                                                                  genome survey devices
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Pred. No.
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; ORGANISM: Homo sapiens
US-09-833-245-1866
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
CT-US01-11988-1866
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Best Local Similarity /2.,
----hes 8; Conservative
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Best Local Similarity 55.
Matches 10; Conservative
                                                                          SEQ ID NO 1866
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                                                                                         NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn Ve
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                                                                                                                              PRIOR FILING DATE:
                                                                                                                                                PRIOR APPLICATION NUMBER:
                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION UNMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
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                                                      LENGTH: 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RIOR APPLICATION NUMBER: 60/199, 384
                                                                                                                                                                                                                                                                                ILE REFERENCE: PF546PCT
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RRENT FILING DATE: 2001-04-12
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                                                                                                                                                60/199; 384
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Pred. No. 30;
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Pred. No. 1
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SEQ ID NO 1864
LENGTH: 63
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Best Local Similarity 55.68
Matches 10; Conservative
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Best Local Similarity
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Sequence 1864, A
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PRIOR FILING DATE: 2000-04-25
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CURRENT EILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILLING DATE: 2000-04-12
PRIOR FILLING DATE: 2000-04-12
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SOFTWARE: Patentin Ve
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PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
                                                                                                                                                                                                                                                                                                                                                                                                      ENERAL INFORMATION
                         NAME/KEY: SITE LOCATION: (37)
                                                                            ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: SITE
LOCATION: (37)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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                                                               FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: PCT/US01/11988
CURRENT FILING DATE: 2001-01-12
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TLE OF INVENTION: Albumin Fusion Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 LCQRPGLVLALPPQLSFS 20
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INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic ; OTHER INFORMATION: at any location having Xaa, Xaa means any amino acid US-09-270-849B-186298
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SEQ ID NO 8
LENGTH: 471
TYPE: PRT
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Best Local Similarity
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'LENGTH: 370
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Best Local Similarity
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NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/270,849B
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                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                          ITLE OF INVENTION: Insect genome survey devices
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PPLICANT: Schulein, Martin
PPLICANT: Outtrup, Helle
PPLICANT: Outtrup, Helle
PPLICANT: Ovel Rhamnogalacturonan Hydrolases
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RPNLXLYYS 251
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Pred. No.
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Pred. No. 3.6e+02;
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Pred. No.
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SEQ ID NO 9884
LENGTH: 393
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LENGTH: 391
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PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
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APPLICANT: - Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
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CURRENT FILING DATE: 2000-12-19
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TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
                             APPLICANT: STAPLETON, JACK T.
TITLE OF INVENTION: FULL-LENGTH GB VIRUS C (HEPATITIS
TITLE OF INVENTION: ARE INFECTIOUS IN PRIMARY CD4 PO
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APPLICATION NUMBER: US/09/828,498
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SCHMIDT, WARREN
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3.9e+02;
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RESULT 13
US-09-724-475-3
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; TYPE: PRT
; ORGANISM: Hepatitis G virus
US-09-828-498-2
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SEQ ID NO 11020
LENGTH: 291
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Best Local Similarity
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APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences
FILE REFERENCE: 38-10(15490)C
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NUMBER OF SEQ ID NOS: 2
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                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                      TITLE OF INVENTION: Recombinant Proteins Of
                                                                                                                                                                                                           NUMBER OF SEQUENCES:
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COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-SOFTWARE: WORDPERFECT 5.1
                                                                                                                      CITY: NEW YORK
STATE: NEW YORK
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                                                                                                                                                            STREET:
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                                                                                                                                                                                                                                                                                          Sergei. A., Emerson,
                                                                                                                                                                                                                                    A Pakistani Strain Of Hepatitis E And Their
                                                                                                                                                                                                                      Use In Diagnostic Methods And Vaccines
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                 PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 32.5; DB 5; Pred. No. 3.5e+02;
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Pred. No. 2.9e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 2910
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; SEQUENCE DESCRIPTION: SEQ ID NO: US-09-724-475-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-769-066-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21, Application US/09769066 GENERAL INFORMATION:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (212) 751-6
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          83 RPGLDLVFA 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 RPGLQLYFS 12
                                 REFERENCE/DOCKET NUMBER: 4600-0293.30
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, v
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/769,066
FILING DATE: 24-Jan-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/542,634
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                        ZIP: 94306
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Dehlinger & Associates
STREET: 350 Cambridge Ave., Suite
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: HEPATITIS E VIRUS ANTIGENS AND USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Fuerst, Thomas R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (212) 758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/947,263 FILING DATE: 18-SEP-1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/724,475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 123 amino acid residues TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: <Unknown>
APPLICATION NUMBER: US08/316,765
FILING DATE: 03-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 28-Nov-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 2026-4032US4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 36,459
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Richard W. Bork
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 08/809,523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     McAtee, C. Patrick
Yarbough, Patrice O.
Zhang, Yifan
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                   324-0960
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 32; DB 5;
Pred. No. 1.8e+02
                                                                                                                                                                                                                                                                                    Version #1.25
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Search completed: June 4, 2001, 12:21:29 Tob time: 591 sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
US-09-270-849B-191061
                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-270-849B-191061
                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 191061
LENGTH: 141
TYPE: PRT
                                                                                                                                                              Query Match
Best Local Similarity 75.0
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 191061, Application US/09270849B
GENERAL INFORMATION:
APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 52.5%; Score 32; DB 5; Best Local Similarity 66.7%; Pred. No. 1.8e+02; Matches 6; Conservative 1; Mismatches 2
                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 RPGLQLYFS 12
|||| | |:
83 RPGLDLVFA 91
                                                                                        1 LLQRPGLQ 8
|:|||| |
66 LIQRPGSQ 73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 124 antho acids
TYPE: amino acid
STRANDEDNESS: Hepatitis E Virus (Burma Strain)
ORF-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                    52.5%;
75.0%;

    Mismatches

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Pred. No. 2.1e+02;
                                                                                                                                                                                                     DB 5; Length 141;
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Database
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64
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                                                                                                                                                               Length
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                1 US-09-702-021-14
PCT-US00-05882-1128
US-08-332-562-94
US-08-462-973-94
US-07-861-895-25
                                                                                                   US-60-160-209-3193
US-60-160-209-4252
US-60-160-203-4414
                                                                                                                                                                                                      SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             _COMB.pep:
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Sequence 3193, Ap
Sequence 4252, Ap
Sequence 4414, Ap
Sequence 7, Appli
Sequence 114, Appl
Sequence 1128, Ap
Sequence 94, Appl
Sequence 94, Appl
Sequence 25, Appl
Sequence 323, Appl
                                                                                                                                                            Description
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-60-	-207-216-74	-60-185-361-50	-60-188-162-350	-60-177-646-232	-09-417-507-425	-60-170-430-24	-09-417-507-365	-60-170-430-262	-60-164-762-67	-60-196-190-107	-09-417-507-2464	-09-417-507-394	-60-155-811-89	-60-155-811-	-60-155-811-56	-60-191-681-1802	37-22	-60-173-464-1873	-60-191-681-1803	-60-191-681-1800	0-191-637-2288	-60-191-637-2283	-60-173-464-1875	-173-464-1871	-60-167-217-2317	-60-167-217-2311	0-167-217-2314	-09-270-767-4546	-60-167-217-2315	T-US01-01324-21	S-09-248-796-26	15-60-245-201-234	-09-417-507-39	-09-634-238-32
e 722,	Φ	equence 508,	equence 3505,	ര	equence	e 2462,	equence 36567,	equence 2622, A	nce 674, Ap	equence	e 2464	equence 39489	e 893,	equence	561,	equence 18020,	æ	equence 1873	e 1803	e 1800	e 2288	equence 2283	equence 18758	equence 18719,	equence 23174,	e 2311	equence 23147,	e 4546	e 23150,		equence 26957,	equence 234, Ap	e 39462,	Ą

## ALIGNMENTS

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; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FASTSEQ FOR W
; SEQ ID NO 3193
; LENGTH: 58
; TYPE: PRT
; ORGANISM: HUMAN
US-60-160-209-3193
US-60-160-209-4252; Sequence 4252, Application US/60160209; GENERAL INFORMATION:
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                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches: 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: BONAZZI, VIVIEN
APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL000113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3193,
                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/60/160,209
CURRENT FILING DATE: 1999-10-19
                                                                                                                     1 FYMGSKTLRGRN 12
|||||||||
25 FYMGSKTLRGRN 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Application US/60160209
                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                         for Windows Version 4.0
                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                   Score 64; DB 23;
Pred. No. 7.5e-05;
Mismatches 0;
                                                                                                                                                                                                                                           Length 58;
                                                                                                                                                                                                       0;
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APPLICANT: Hogarth, P. Mark
APPLICANT: Powell, Maree S.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: Maxwell, Kelly F.
APPLICANT: Garrett, Thomas P.J.
APPLICANT: Epa, Vidana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ORGANISM: HUMAN
US-60-160-203-4414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             B
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Thes 12; Conservat
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Matches 12; Conservative C
                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ
SEQ ID NO 4414
LENGTH: 102
                                                          TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -60-160-203-4414
Sequence 4414, Application US/60160203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/60/160,203
CURRENT FILING DATE: 1999-10-19
NUMBER OF SEQ ID NOS: 6374
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OTHER INFORMATION: X
-60-160-209-4252
                                              [LE REFERENCE: 4102-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENERAL INFORMAT
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CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TILE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
ITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
ITLE OF INVENTION: USES THEREOF
IT BEFORE COLORS OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF THE SECRET OF 
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FITLE OF INVENTION: ISOLATE
FITLE OF INVENTION: ACID MODIFIED OF INVENTION: THEREOF
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                                                                                                                                                                                                                                                                                                                                                                                                              Application US/09245764
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ON NUMBER: US/09/245,764
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          THEREOF
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Pred. No. 0.00014;
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Pred. No.
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; TYPE: PRT
; ORGANISM: HOMO Sapiens
US-09-245-764-7
                                                                                                                          SOFTWARE: PatentIn
SEQ ID NO 1128
LENGTH: 399
                                                                                                                                                                                                                                                                                                                    Sequence 1128, Application GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                   CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124/270
EARLIER FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
FEATURE:
NAME, KEY: SITE
LOCATION: (208)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 14
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Best Local Similarity
Matches 12; Conserv
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EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                 APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
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                                                                                                                                                                                                                                                   ITLE OF INVENTION: Human Cancer Associated ILE REFERENCE: PA106PCT
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EARLIER FILING DATE: 1998-09-11
EARLIER APPLICATION NUMBER: 60/073,972
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Pred. No.
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Pred. No. 0.00039;
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                                                                                                                                                                                                                                                          Sequences and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 374;
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; LOCATION: (349) ; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids PCT-US00-05882-1128

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Best Local Similarity 100.
Matches 12; Conservative
                                                                                Query Match
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                                                                                                                                                                                                            FORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                             CLASSIFACE CONTROL DATA:
PRIOR APPLICATION NUMBER: US 07/896,457
PATTING DATE: 27-MAY-1992
                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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254 FYMGSKTLRGRN 265
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1 FYMGSKTLRGRN 12
||:||| | ||
41 FYVGSKILEYRN 52
                                                                      Local Similarity
                                                                                                                                                                                                                                                                                              NAME: BENT, Stephen A REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                           TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/0 FILING DATE: 31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
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                                                                                                                                                                                  LENGTH:
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                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HULETT, Mark D.
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                                                                                                                                           linear
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, Mark P.
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                                                                                                                                                                                                                                                                                             29,768
                                                                                                                                                                                                              94:
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                                                                      Score 40;
Pred. No.
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                                                       Mismatches
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                                                                                   DB 7;
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                                                                                 Length 91;
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                                                       Indels
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                                                       Gaps
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RESULT 8 US-08-462-973-94

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US-07-861-895-25

GENERAL INFORMATION:

Application US/07861895

HOGARTH, Phillip M. HULETT, Mark D.

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Query Match
Best Local Similarity
Conservi
                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR OF TOWNS ON NUMBER: US 08/332,562
                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: AU PM8232/94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UMBER OF SEQUENCES:
4
                                                                                                                                                                         LENGTH: 91 amino acids TYPE: amino acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: WO PCT/AU87/00159 FILING DATE: 29-MAY-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
                                                                                                                                                                                                                                                                     TELEFAX:
                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                      NAME: BENT, Stephen A. REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 20007-5109
                              1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                      PPLICATION NUMBER:
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FYVGSKILEYRN 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/08462973
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                                                             Conservative
                                                                                                                                                                                                                                                                     (202)672-5399
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HULETT, Mark D.
MENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HOGARTH, Mark P.
HIBBS, Margaret L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MCKENZIE, Ian F.C.
                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                       (202)672-5300
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26-MAY-1986
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                                                                                                                                                                                                                                                                                                                         16786/203
                                                                         Score 40; DB
Pred. No. 5.4
                                                               Mismatches
                                                                                             DB 8;
                                                                                        Length 91;
                                                             Indels
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                                                               Gaps
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IERINO, Francesco

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Query Match
Best Local Similarity ou.,
Thes 8; Conservative
                                 FILE REFERENCE: 11000.1043U1
CURRENT APPLICATION NUMBER: U
CURRENT FILING DATE: 2000-08
                                                                       APPLICANT: Reid, Julian R.
APPLICANT: Coolbear, Timothy
TITLE OF INVENTION: Polynucleotides,
TITLE OF INVENTION: them and methods
                                                               TILE REFERENCE:
                                                                                                                                                                                                                                                                                            09-634-238-323
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FILING DATE: 25-OCT-1989
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,76
REFERENCE/DOCKET NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 410 amino acid
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein -861-895-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25
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TOPOLOGY: linear

LECUTO TO TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 25-OCT-1
PRIOR APPLICATION DATA:
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COMPUTER: IBM PC COMPATIBLE

COPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
                                                                                                                                                                                                                                                               INFORMATION:
      SEQ ID NOS:
FastSEQ fo.
                                                                                                                                     Holland, Ross
O'Toole, Paul W.
                                                                                                                                                                   Christensson, Anna C.
                                                                                                                                                                                 Lubbers, Mark
Dekker, James
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Bloksberg, Leonard, N.
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                                                                                                                                                                                                                                                                         Application US/09634238
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3000 K Street, N.W.,
ashington, D.C.
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(202)672-5399
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for Windows Version
                           NUMBER: US/09/634,238
2000-08-08
                                                                                                                                                                                                 Mark W.
                                                                                                                                                                                                                                                                                                                                                                                                                         62.5%;
66.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                        Score 40;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                    materials incorporating for using them.
 4.0
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Query Match
Best Local Similarity

57.8%; 45.5%;

Score 37; Pred. No.

DB 18;

Length 197;

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; ORGANISM: A.fumigatus
US-09-417-507-39462
                                                                                                                                                                                        US-09-417-507-39462
; Sequence 39462, Application US/09417507
                                                                                                                                                                                                                                                                       B
                               SEQ ID NO 39462
LENGTH: 197
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Lactobacillus rhamnosus US-09-634-238-322
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Best Local Similarity
7; Conserv
                                                                                                                                                               GENERAL INFORMATION: APPLICANT: KEITH G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Lactobacillus rhamnosus US-09-634-238-323
                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 322
                                                                                       CURRENT APPLICATION NUMBER: US/09/417,507 CURRENT FILING DATE: 1999-10-14
                                                                                                                      APPLICANT: KEITH G. WEINSTOCK ET AL.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ASPERGILLUS
TITLE OF INVENTION: FUMIGATUS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                         NUMBER OF SEQ ID NOS:
                                                                                                                   FILE REFERENCE: PATH99-10
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LENGTH: 268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE:
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                                                                                                                                                                                                                                                                  407 YVGQKPLRSRN 417
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                                                                                                                                                                                                                                                                                              2 YMGSKTLRGRN 12
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                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FastSEQ for Windows Version 4.0
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Bloksberg, Leonard,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Coolbear, Timothy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reid, Julian R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     O'Toole, Paul W.
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Holland, Ross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dekker, James
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jubbers, Mark W.
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                                                                                                                                                                                                                                                                                                                                         59.4%;
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Pred. No.
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Pred. No. 43;
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1.3e+02;
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                                                                                                                                                                                                                                                                                                                    Gaps
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Query Match
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US-09-248-796-26957
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US-09-248-796-26957
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US-60-245-201-234
                                                                                        Sequence 2181, Application PC/TUS0101324
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26957, Appl
GENERAL INFORMATION:
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LENGTH: 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 1999-02-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/248,796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PPLICANT: Keith Weinstock et al PPLICANT: Keith Weinstock et al IITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAIITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
PILE REFERENCE: 107196.132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/60/245,201
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                PPLICANT: Human Genome Sciences, Inc., et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, a
TLE REFERENCE: PC002PCT
URRENT APPLICATION NUMBER: PCT/US01/01324
URRENT FILING DATE: 2001-01-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 1845
rior application data removed - refer to PALM or file wrapper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
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89 FKIGSKTLEGQ 99
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DRUG-METABOLIZING PROTEINS,
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66.7%;
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                                                            Best Local Similarity 87. Matches 7; Conservative
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2181
LENGTH: 153
                                                                                                                                                               OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                                                                                                                                 NAME/KEY: SITE LOCATION: (70)
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OTHER INFORMATION:
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Search completed: June 4, 2001, 12:20:03 Job time: 516 sec

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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2: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*

3: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

4: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

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 Query Match
Best Local Similarity 100
Matches 12; Conservative

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Score 64; D Pred. No. 4. Mismatches

DB 5; 1.7e-06;

Length 12; ; ; Indels

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US-09-459-573-12

Sequence 12, Application US/09459573 GENERAL INFORMATION:

APPLICANT: LIVSHITS, VITALIY
APPLICANT: ZAKATAEVA, NATALIA
APPLICANT: NAKANISHI, KAZUO
APPLICANT: VENIANINOVICH, VLADIMIR

APPLICANT: TOKHMAKOVA, IRINA
TITLE OF INVENTION: METHOD FOR PRODUCING L-AMINO ACIDS
FILE REFERENCE: 0010-1066-0
CURRENT APPLICATION NUMBER: US/09/459,573
CURRENT FILING DATE: 1999-12-13

APPLICANT:

TROSHIN, PETR

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ORGANISM: Conus radiatus CT-US00-01979A-73
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Sequence 69, Application PC/TUS0001979A

GENERAL INFORMATION:
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                                                        SEQ ID NO 73
LENGTH: 61
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SEQ ID NO 69
LENGTH: 61
                                                                                     NUMBER OF SEQ ID NOS: 404
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                              FILE REFERENCE: Alphas 2
CURRENT APPLICATION NUMBER: PCT/US00/01979A
CURRENT FILING DATE: 2000-01-28
                                                                                                                          PRIOR APPLICATION NUMBER: US 60/118/381
PRIOR FILING DATE: 1999-01-29
                                                                                                                                                                                                                                        equence 73, Application PC/TUS0001979A
BNERAL INFORMATION:
APPLICANT: University of Utah Research Foundation
APPLICANT: Cognetix, Inc.
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Best Local Similarity
Matches 8; Conserv
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CURRENT FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: US 60/118,381
PRIOR FILING DATE: 1999-01-29
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TITLE OF INVENTION: Alpha-Conotoxin Peptides
FILE REFERENCE: Alphas 2
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NUMBER OF SEQ ID NOS: 24
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                                                                                                                                                                                                          LE OF INVENTION: Alpha-Conotoxin Peptides
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17 FTSGRSTFRGRN 28

FYMGSKTLRGRN 12

Query Match

Local

Similarity 7; Conserv

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Score 35; DB Pred. No. 11;

Length 61;

Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Conus textile US-09-493-795A-69
                               SEQ ID NO 73
LENGTH: 61
TYPE: PRT
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LENGTH: 61
TYPE: PRT
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Best Local Similarity . 58.3%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                          sequence /3,
                                                                          NUMBER OF SEQ ID NOS: 404
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                         PRIOR APPLICATION NUMBER: US 60/118,381 PRIOR FILING DATE: 1999-01-29
                                                                                                                                  TITLE OF INVENTION: Alpha-Conotoxin Peptides FILE REFERENCE: 2314-179.A CURRENT APPLICATION NUMBER: US/09/493,795A CURRENT FILING DATE: 2000-01-28
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Best Local (
            ORGANISM: Conus radiatus
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                                                                                                                                                                                                                                                                     NERAL INFORMATION:
PPLICANT: Watkins, Maren
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Hillyard, David R.
McIntosh, J. Michael
                                                                                                                                                                                                                                                                                                      Application US/09493795A
                                                                                                                                                                                                         Jones, Robert M.
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McIntosh, J. Michael
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58.3%;
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CT-US00-01979A-65

equence 65, Application PC/TUS0001979A

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Best Local Similarity 50...
7; Conservative
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Best Local
                                               GENERAL INFORMATION:
                                                           Sequence 12089, Application US/09739449
                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PA
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SEQ ID NO 65
LENGTH: 64
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NUMBER OF SEQ ID NOS: 404
                                 APPLICANT: Hinkle, Gregory J.
                                                                                                                                                                                                                                                                                                          LENGTH: 64
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/493,795A
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                                                                                                                                                                                                                                                                                          ORGANISM: Conus textile
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APPLICATION NUMBER: US 60/118,381
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                                                                                                                                         FSSGRSTFRGRN 28
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INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
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                 Slater,
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Query Match Best Local Similarity

51.68;

Score 33; Pred. No.

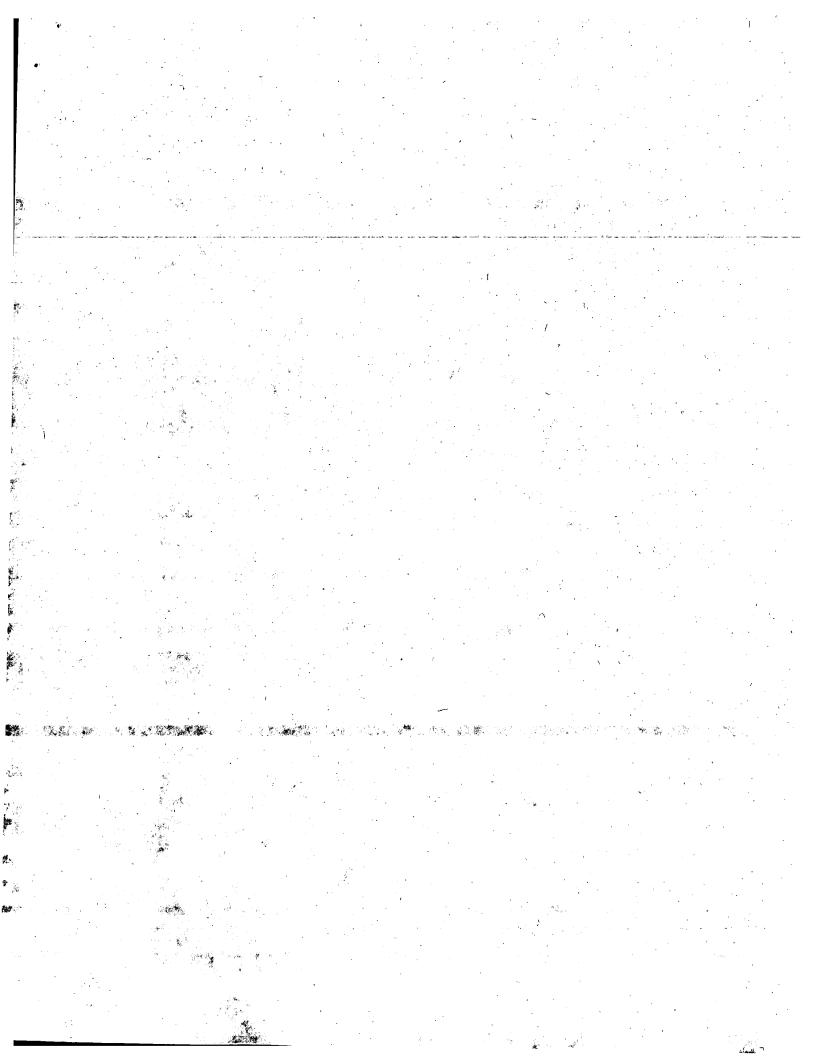
DB 5; 24;

Length 55;

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; FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: at any location having Xaa, Xaa means any amir US-09-270-849B-194148
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Best Local Similarity
Thehes 6; Conserv
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US-09-270-849B-194148
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                                                                                               SOFTWARE: PatentIn Ver.
SEQ ID NO 194148
LENGTH: 55
TYPE: PRT
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SEQ ID NO 12004
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LENGTH: 377
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Best Local Similarity
                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17.
NUMBER OF SEQ ID NOS: 195450
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CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                 APPLICANT: Swimmer et al TITLE OF INVENTION: Insec
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TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15490)C
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TYPE: PRT
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                                                                           ORGANISM: Artificial Sequence
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Pred. No.
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Query Match
Best Local Similarity
6; Conserv
                                                          ORGANISM: Homo sapiens
CT-US01-11988-458
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; OTHER INFORMATION: Incyte ID No: 2879792
US-09-823-356-16
           Query Match
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                                                                                                                                     SEQ. ID NO 458
                                                                                                                                                 NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn Ver.
                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 60)
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60,
PRIOR FILING DATE: 2000-04-25
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LENGTH: 275
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PRIOR FILING DATE: 2000-04-1
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                                                                                                                 ENGTH:
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URRENT FILING DATE: 2001-01-12
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SOFTWARE: PERL Program
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Baughn, Mariah R.
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 Score 33; DB 1;
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Pred. No. 1.3e+02;
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Length 282;
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Best Local Similarity 93...
Change 6; Conservative
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US-09-833-245-458
                                                      PRIOR APPLICATION NUMBER: US 09/440,676 PRIOR FILING DATE: 1999-11-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 458
LENGTH: 282
                                   PRIOR APPLICATION NUMBER: US 09/440,677
                                                                                                     PRIOR FILING DATE: 1999-11-15
PRIOR APPLICATION NUMBER: US 09/520,478
                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/440,493
                                                                                                                                                              PRIOR APPLICATION NUMBER: US 09/440,370 PRIOR FILING DATE: 1999-11-12
                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 09/439,878
                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 09/268,865
PRIOR FILING DATE: 1999-03-15
PRIOR APPLICATION NUMBER: US 09/450,810
PRIOR ETLING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: US 09/453,137
PRIOR APPLICATION NUMBER: US 09/453,137
PRIOR APPLICATION NUMBER: US 09/453,137
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Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/525,361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                        TTLE OF INVENTION: NOVEL METHODS OF DIAGNOSING AND TITLE OF INVENTION: COMPOSITIONS, AND METHODS OF SCITLE OF INVENTION: CANCER MODULATORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/833,245
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ENERAL INFORMATION:
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PPLICANT: GISH, KURT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TERAL /INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         LE REFERENCE: A-67860-3/DJB/JJD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            46 YLGSKTL 52
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                      1999-11-16
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85.7%;
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l; Mismatches 0;
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                                                                            100.0
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Gapop 10.0 , Gapext 0.5
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57
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                                                                                                                                                                                       Length
                                                                                                      US-60-160-209-3193
US-60-160-209-4252
US-60-160-203-4414
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Sequence 3193, Ap
Sequence 4252, Ap
Sequence 4214, Ap
Sequence 7, Appli
Sequence 14, Appl
Sequence 1128, Ap
Sequence 1129, Ap
Sequence 621, App
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149,	21673,	e 5274	,	152, 4	615, 7	167,	78	78		6400	Sequence 932, App	1490	Sequence 12472, A	37264,	Sequence 4715, Ap	6123,	e 7854,	e 6355,	e 7951, A	3644	2559	e e	253	۵	2	2	e 2,	Sequence 16876, A	2		4	4,	8221, 4	•

## ALIGNMENTS

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; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSEQ for W
; SEQ ID NO 3193
; LENGTH: 58
; TYPE: PRT
; ORGANISM: HUMAN
US-60-160-209-3193
US-60-160-209-4252

; Sequence 4252, Application US/60160209

; GENERAL INFORMATION:
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Best Local Similarity
Matches 12; Conser
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CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID. MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CL000113
                                                                                                                       37 TSSEYQILTARR 48
                                                                                                                                                             1 TSSEYQILTARR 12
                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/60160209
                                                                                                                                                                                                                                                                                                                                                                                                       for Windows Version 4.0
                                                                                                                                                                                                                     100.0%; Score 57; DB 23; 100.0%; Pred. No. 0.00023
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                                                                                                                                                                                                                                          Length 58;
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: LOCATION: (1)...(96)
: OTHER INFORMATION: Xaa = Any Amino Acid
US-60-160-209-4252
                                                                                                                                                                                                     Sequence 7, Application US/09245764 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           Query Match 100
Best Local Similarity 100
Matches 12; Conservative
                         APPLICANT: Hogarth, P. Mark
APPLICANT: Powell, Maree S.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: McKenzie, Ian F.C.
APPLICANT: Maxwell, Kelly F.
APPLICANT: Maxwell, Kelly F.
APPLICANT: Epa, Vidana
TITLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF FC RECEPTORS
TITLE OF INVENTION: AND USES THEREOF
FILE REFERENCE: 4102-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 6374

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4414
LENGTH: 102
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Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                               -09-245-764-7
CURRENT APPLICATION NUMBER: US/09/245,764
CURRENT FILING DATE: 1999-02-05
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LENGTH: 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
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CURRENT FILING DATE: 1999-10-19
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                                                                                                                                                                                                                                                                                                                      63 TSSEYQILTARR 74
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                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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KINASE PROTIENS AND USES
                                                                                                                                                                                                                                                                                                                                                                                                                             Length 102;
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; SEQ ID NO 7; LENGTH: 261; TYPE: PRT; ORGANISM: Homo sapiens US-09-245-764-7
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                                                                                                                            CURRENT APPLICATION NUMBER: PCT/US00/05882
CURRENT FILING DATE: 2000-03-08
EARLIER APPLICATION NUMBER: 60/124,270
EARLIER FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1128
LENGTH: 399
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                                                                                                                                                                                                                                                                                                                                    Sequence 1128, Application PC/TUS0005882 GENERAL INFORMATION:
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                  APPLICANT: Craig Rosen,
APPLICANT: Steve Ruben
TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: C: Fras
TITLE OF INVENTION:
FILE REFERENCE: 78
                              NAME/KEY: SITE LOCATION: (208) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EARLIER APPLICATION NUMBER: 60/099,994
EARLIER FILING DATE: 1998-09-11
EARLIER APPLICATION NUMBER: 60/073,972
EARLIER FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 15
                                                                                             ORGANISM: Homo sapiens
                                                                                                                 TYPE: PRI
                                                                               FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
WAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                     241 TSSEYQILTARR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           222 TSSEYQILTARR 233
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mes 12; Conserv
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                       Xaa equals any of the naturally occurring L-amino acids
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Pred. No.
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Pred. No.
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; LOCATION: (349) ; OTHER INFORMATION: Xaa equals any PCT-USO0-05882-1128

of the naturally occurring L-amino

acids

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US-60-181-426-621
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                                                                                                                                                                            SEQ ID NO 621
LENGTH: 56
                                                                                                                                                                                                                                                                                                                               Sequence 621, Application U GENERAL INFORMATION:
APPLICANT: Bonazzi, Vivien
                                                            Matches
                                                                          Query'Match
Best Local 9
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Best Local
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SOFTWARE: FastSEQ for
SEQ ID NO 1209
LENGTH: 56
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Best Local Similarity
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ORGANISM: HUMAN
-60-173-468-1209
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                                                                                                                                                                                                             CURRENT FILING DATE:
NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/60/181,426
CURRENT FILING DATE: 2000-02-09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             URRENT APPLICATION NUMBER: US/60/173,468
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ITLE OF INVENTION:
ITLE OF INVENTION:
                                                                                                                                                                   TYPE: PRT
                                                                                                                                                                                                                                                                                   ITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS, ITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS, ITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PPLICANT: Bonazzi,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
ses 7; Conserv
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                             2 SSEYQILTAR 11
SSEYETLTAK 24
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                                                                            Similarity
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                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AND USES THEREOF
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                                                                          63.2%;
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                                                                         Score 36; Pred. No.
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Pred. No.
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                                                                                     Length 56;
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                                                           Indels
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RESULT

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
FILING DATE: 27-MAY-1992
ATTORNEY/AGENT INFORMATION:
                                             CORRESPONDENCE ADDRESS:
                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                          53 TSSEYHIARAER 64
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                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                      1 TSSEYQILTARR 12
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INVENTION: FC REC
Washington
                                                                          INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                                                Application US/08462973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08332562
              3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3000 K Street, N.W
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HIBBS, Margaret L.
                                                                                                        BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                    HIBBS, Margaret L
                                                                                                                                                                  MCKENZIE,
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                                                                                        HULETT, Mark D.
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                             Foley
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                                                                                                                       Bernadette M.
                             & Lardner
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                                                                                                                                                     Mark
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                                                                                                                                                                                                                                                                                                                                                    Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Suite 500
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                                                                                                                                                                                                                                                                                                                                                               Length 91;
                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                    Gaps
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OMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

PatentIn Release #1.0, Version #1.30

ZIP: 20007-5109

Best Local Similarity 70.0 Matches 7; Conservative

70.0%;

Pred. No. 17; 2; Mismatches

Indels

0; Gaps

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JS-60-207-216-686
                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity bu...
Ahes 8; Conservative
                                                               CURRENT APPLICATION NUMBER: US/60/207,216
CURRENT FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 898
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 686
                                                                                                                                             APPLICANT: Beasley,
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: CLO
ORGANISM: HUMAN
-60-207-216-686
                                                                                                                                                                                                                                                      Sequence 686,
                               LENGTH: 15
TYPE: PRT
                                                                                                                                                                                                                                   ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
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APPLICATION NUMBER: WO PCT/AU87/00159
FILING DATE: 29-MAY-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
PTI-ING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/332,562
PILING DATE: 31-OCT-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
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                                                   151
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                                                                                                                                                                                                                   Beasley, Ellen
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linear
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26-MAY-1986
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                                                                                                                                                                  AND USES THEREOF
                                                                               Windows Version 4.0
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                                                                                                                                                                           ENCODING HUMAN DRUG TARGET PROTEINS,
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Query Match

63.2%;

Score 36; DB 23;

Length 151;

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RESULT 14
US-09-718-815-4
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Best Local Similarity
7; Conserv
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                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                            Sequence 4,
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Best Local Similarity 50...
Thes 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/718,692
CURRENT FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/489,039
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Human
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IITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
IITLE OF INVENTION: PREUMONIAE FOR DIAGNOSTICS A
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TYPE: PRT
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254 SSEYETLTAK 263
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                                        INVENTION:
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                                                                                                  Beraud, Christophe
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VENTION: Novel motor proteins and methods for VENTION: their use
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, Richard
Novel motor proteins and methods for
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58.3%;
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Pred. No. 42;
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Sequence 4, Application US/09718852

GENERAL INFORMATION:
APPLICANT: Beraud, Christophe
APPLICANT: Ereedman, Richard
TITLE OF INVENTION: Novel motor proteins and methods for
TITLE OF INVENTION: their use
FILE REFERENCE: 1052
CURRENT APPLICATION NUMBER: US/09/718,852
CURRENT FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 366
TYPE: PAT
ORGANISM: Human
US-09-718-852-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FITLE OF INVENTION: their use
FILE REFERENCE: 1052
CURRENT APPLICATION NUMBER: US/09/718,815
CURRENT FILING DATE: 2000-11-22
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 366
TYPE: PRT
ORGANISM: Human
US-09-718-815-4
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US-09-718-852-4
                                                                       Query Match 63.:
Best Local Similarity 70.0
Matches 7: Conservative
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Best Local Similarity 70.0%;
Matches 7; Conservative
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254 SSEYETLTAK 263
2 SSEYQILTAR 11
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254 SSEYETLTAK 263
                                                                                            63.2%; Score 36; DB 21; 70.0%; Pred. No. 48;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Match
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2: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:*

3: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

4: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*

5: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*

6: /cgn2_6/ptodata/1/paa/US09_NEW_COMB.pep:*
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US-09-770-849B-180859
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US-09-779-449-12004
US-09-739-449-8153
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PCT-US01-11988-763
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US-09-739-449-11547
US-60-248-823-102
US-09-739-449-8297
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PCT-US00-35017A-1460
PCT-US01-06769-11
PCT-US01-11988-762
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PCT-US01-04098A-1753
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Sequence 1460, Ap
Sequence 11, Appl
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721, App
1872, Ap
1872, Ap
1800, Ap
180859,
190000,
12004, A
8153, Ap
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763, App
11547, A
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1753, Ap
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		Sequence 6, Appli	Sequence 4, Appli	Sequence 1566, Ap	Sequence 8964, Ap	Sequence 1035, Ap	Sequence 3003, Ap	Sequence 581, App	Sequence 3665, Ap	Sequence 1697, Ap	Sequence 11488, A	Sequence 188275,	Sequence 8568, Ap	Sequence 991, App	Sequence 1236, Ap	Sequence 89, Appl	Sequence 2, Appli	Sequence 4, Appli	Sequence 82, Appl	

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RESULT 2 PCT-US00-35017A-1460 Sequence 1460, Application PC/TUS0035017A GENERAL INFORMATION: APPLICANT: Hyseq Inc TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides FILE REFERENCE: 784PCT CURRENT APPLICATION NUMBER: PCT/US00/35017A CURRENT FILLNG DATE: 2000-12-2 PRIOR APPLICATION NUMBER: US09/488,725 PRIOR FILING DATE: 2000-01-21 PRIOR FILING DATE: 2000-01-21 PRIOR FILING DATE: 2000-01-21 PRIOR FILING DATE: 2000-01-25 PRIOR FILING DATE: 2000-04-25 NUMBER OF SEQ ID NOS: 1478	Oy 1 TSSEYQILTARR 12	Query Match 100.0%; Score 57; DB 5; Length 12; Best Local Similarity 100.0%; Pred. No. 1.9e-05;Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps	OTHER INFORMATION: Oligopeptides representing the extracellular; OTHER INFORMATION: domain of CD64 US-09-284-107-25	25 12 T T : Artificial Sequence	ובי יכו	CURRENT APPLICATION NUMBER: US/09/284,107  PRIOR APPLICATION NUMBER: PCT/NL97/00557  PRIOR FILING DATE: 1997-10-07	; APPLICANT: Logtenberg, Ton ; APPLICANT: de Kruif, Cornelis Adriaan John ; TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES ; TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FOR A TARGET ; TITLE OF INVENTION: 313633000600	RESULT 1 US-09-284-107-25 ; Sequence 25, Application US/09284107 ; GENERAL INFORMATION:

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Query Match
Best Local Similarity
5; Conserve
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Best Local Similarity
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PCT-US00-35017A-1460
                                                                                                                                             SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 762
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PRIOR FILING DATE: 2000-03-02
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 11
OTHER INFORMATION: Xaa equals -US01-11988-762
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PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
                                                                                                                                                                                NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                               PRIOR FILING DATE:
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CURRENT FILING DATE: 2001-03-02
                                                                             TYPE: PRT
ORGANISM: Homo sapiens
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TITLE OF INVENTION: hB7;H2, A NOVEL CO-STIMULATORY MOLECULE
FILE REFERENCE: 07039-202WO1
                                                  NAME/KEY: SITE
                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                      URRENT APPLICATION NUMBER: PCT/US01/11988
URRENT FILING DATE: 2001-01-12
                                                                                                                             ENGTH:
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ITLE OF INVENTION: Albumin Fusion Proteins
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                LE REFERENCE: PF546PCT
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   any of the naturally occurring L-amino acids
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Pred. No. 70;
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Query Match
Best Local Similarity 45...
"hes 5; Conservative
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; Sequence 763, App
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; LOCATION: (110)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-833-245-762
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SEQ ID NO 763
LENGTH: 290
TYPE: PRT
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Best Local Similarity 43...
5; Conservative
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Best Local Similarity 45.3
"""" 65; Conservative
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PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
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Sequence 762, Application US/09833245
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PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
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CURRENT FILING DATE: 2001-01-12
CURRENT APPLICATION NUMBER: 60/229, 358
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NUMBER OF SEQ ID NOS: 2267
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PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
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CURRENT FILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229, 358
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                        Score 32; DB 1;
Pred. No. 96;
5; Mismatches
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Pred. No. 80;
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Pred. No. 80;
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US-60-248-823-102; Sequence 102, Application US/60248823; GENERAL INFORMATION:
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Whiches 5; Conserve
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SEQ ID NO 11547
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CURRENT FILING DATE: 2000-12-19
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PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/199, 384
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TILE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
TLE REFERENCE: 38-10(15490)C
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TYPE: PRT
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URRENT FILING DATE: 2001-04-12
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45.5%;
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Pred. No. 1.2e+02;
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Pred. No.
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US-60-248-823-102
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PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 8297
LENCTH: 240
TYPE: PRT
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GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 102
LENGTH: 715
TYPE: PRT
SOFTWARE: Pa
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                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                               TITLE OF INVENTION: Albumin Fusion Proteins FILE REFERENCE: PF546PCT
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APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
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                                                             PRIOR FILING DATE:
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                                                                                                                                                                                           CURRENT APPLICATION NUMBER: PCT/US01/11988
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                                                                             APPLICATION NUMBER: 60/199, 384
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Pred. No. 1
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Pred. No.
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; ORGANISM: Homo sapiens PCT-US01-11988-133

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Query Match
Best Local Similarity
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Sequence 1378, Application PC/TUS0104098A
PRIOR FILING DATE: 2000-09-01
PRIOR APPLICATION NUMBER: 09/620,325
PRIOR FILING DATE: 2000-07-19
PRIOR APPLICATION NUMBER: 09/598,075
PRIOR APPLICATION NUMBER: 09/560,875
PRIOR APPLICATION NUMBER: 09/560,875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver; 2.1
SEQ ID NO 133
                                                                                                                                                                                                        JURRENT APPLICATION NUMBER: PCT/US01/04098A JURRENT FILING DATE: 2001-02-05 PRIOR APPLICATION NUMBER: Not Yet Assigned PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: 09/728,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-833-245-133
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Best Local Similarity
Matches 7; Conserv
                                                                                                                           RIOR APPLICATION NUMBER: 09/663,561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 133;
                                                                                                                                                             RIOR APPLICATION NUMBER: 09/693,325
                                                                                                                                                                                                                                                                                                                            PPLICANT: Hyseq, Inc.
                                                                                                                                                                                                                                                                                                                                                              NERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2001-04-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR APPLICATION NUMBER: 60/256, 931
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Human Genome SCIENCE, Proteins TITLE OF INVENTION: Albumin Fusion Proteins REFERENCE: PF546PCT 'MC/NG/R33.245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                             LE REFERENCE: 21272-029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   222 SSEFQIPTA 230
                                                                                                        APPLICATION NUMBER: 09/654,936
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                                                                                                                                                                                                  FILING DATE: 2000-11-30
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77.88;
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77.88;
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Pred. No. 2e+02;
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Pred. No.
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PCT-US01-04098A-1753
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; ORGANISM: Mus musculus
US-09-193-562D-34
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Best Local Similarity burning 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1753, Application PC/TUS0104098A
                                                                                                                                                                                                                                                                             FILE REFERENCE: 21272-029
CURRENT APPLICATION NUMBER: PCT/US01/04098A
CURRENT FILING DATE: 2001-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 34, Applica GENERAL INFORMATION:
                                                                                                            PRIOR FILING DATE:
                                                                                                                                                       PRIOR APPLICATION NUMBER: 09/663,561
                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 09/728,422
PRIOR FILING DATE: 2000-11-30
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                                                                                                                                                                                PRIOR APPLICATION NUMBER: 09/693,325 PRIOR FILING DATE: 2000-10-20
                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: Not yet Assigned PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides FILE REFERENCE: 21272-029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Pauli, Benédicht U.
TITLE OF INVENTION: Nucleotide Sequences Encoding Mammalian Calcium TITLE OF INVENTION: Activated Chloride Channel-Adhesion Molecules FILE REFERENCE: 18617.0052
CURRENT APPLICATION NUMBER: US/09/193,562D
CURRENT FILING DATE: 1998-11-17
PRIOR APPLICATION NUMBER: US/60/065,922
PRIOR FILING DATE: 1997-11-17
PRIOR FILING DATE: 1997-11-17
NUMBER OF SEQ ID NOS: 47
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Best Local Similarity 62:

Matches 5; Conservative
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SOFTWARE: Custom
SEQ ID NO 1378
LENGTH: 790
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                                  APPLICATION NUMBER: 09/598,075 FILING DATE: 2000-06-20
                                                                    APPLICATION NUMBER: 09/620,325
FILING DATE: 2000-07-19
                                                                                                                   APPLICATION NUMBER: 09/654,936
APPLICATION NUMBER: 09/560,875 FILING DATE: 2000-04-27
                                                                                                                                            FILING DATE: 2000-09-15
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                                                                                                        2000-09-01
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Pred. No. 5.6e+02;
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Pred. No. 4.8e+02;
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PRIOR FILING DATE: 2000-02-03

NUMBER OF SEQ ID NOS: 3960

SOFTWARE: Custom
SEQ ID NO 1753

LENGTH: 4544

TYPE: PRT
CORGANISM: Homo sapiens
PCT-US01-04098A-1753.

Query Match
Best Local Similarity 62.5%; Pred. No. 3.4e+03;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Search completed: June 4, 2001, 12:21:29

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Result No.	Score	Query Match Length DB ID	ength	₽.	ID	Description
, ,	. 57	89.1	96	23	us-60-160-209-4252	Sequence 4252, Ap
2	57	89.1	102	23	US-60-160-203-4414	Sequence 4414, Ap
w	57	89.1	261	16	US-09-245-764-7	Sequence 7, Appli
4	57	89.1	374	21	US-09-702-021-14	Sequence 14, Appl
5	57	89.1	399	ר	PCT-US00-05882-1128	Sequence 1128, Ap
6	51	79.7	81	23	US-60-170-373-3410	Sequence 3410, Ap
, 7	51	79.7	18	23	US-60-177-571-3365	Sequence 3365, Ap
80	50	78.1	93	18	US-09-474-434-1676	Sequence 1676, Ap
9	50	78.1	93	23	US-60-173-463-626	Sequence 626, App
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EDSGLYQCEAAJ             EDSGLYWCEAAJ	lar Con	ORGANISM: HUMAN FEATURE: VARIANT LOCATION: (1)( OTHER INFORMATION 0-160-209-4252	ATI DA ID SEQ	ISULT 1  :60-160-209-4252 Sequence 4252, Application: GENERAL INFORMATION: APPLICANT: BONAZZI, TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
EDSGLYQCEAAT 12	ity ser	N: .	ON NOS fo	ZZZHZI
12 78	h 89. Similarity 91. 11; Conservative	) Xaa	CURRENT APPLICATION NUMBER CURRENT FILING DATE: 1999- NUMBER OF SEQ ID NOS: 4646 SOFFWARE: FASTSEQ for Wind EQ ID NO 4252 LENGTH: 96 TYPE: PRT	ation VIVIEN VISOLAT ACID M
	Query Match Best Local Similarity 91.7% Matches 11; Conservative	<b>1</b> -	CURRENT APPLICATION NUMBER: US/ CURRENT FILING DATE: 1999-10-19 NUMBER OF SEQ ID NOS: 4646 SOFTWARE: FASTSEQ FOR WINDOWS V EQ ID NO 4252 EQ ID NO 4252 TYPE: PRT	SULT 1  :60-160-209-4252  Sequence 4252, Application US/60160209  GENERAL INFORMATION: APPLICANT: BONAZZI, VIVIEN TITLE OF INVENTION: ACID MOLECULES EN  TITLE OF INVENTION: THEREOF
•	18; 78;	Any	: U	OS/
	0;	Am	s/6 19 ve:	HUM.
	Med	· o	0/16 rsic	5020
	Score 57; DB Pred. No. 0.0 ; Mismatches	ORGANISM: HUMAN FEATURE: NAME/KEY: VARIANT LOCATION: (1)(96) OTHER INFORMATION: Xaa - Any Amino Acid O-160-209-4252	CURRENT APPLICATION NUMBER: US/60/160,209 CURRENT FILING DATE: 1999-10-19 NUMBER OF SEQ ID NOS: 4646 SOFTWARE: FASTSEQ for Windows Version 4.0 EQ ID NO 4252 LENGTH: 96 TYPE: PRT	)9 (INA
	); o. c	<u>.</u>	209	NIDIN
	Score 57; DB 23; Pred. No. 0.015; Mismatches	ě		ation US/60160209 VIVIEN ISOLATED HUMAN KINASE PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN KINASE PR
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RESULT 4
US-09-702-021-14
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                                  Sequence 14, Application US/09702021
GENERAL INFORMATION:
APPLICANT: C. Fraser
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Best Local 9
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF FILE REFERENCE: 7853-217
                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 7
LENGTH: 261
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Best Local Similarity
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SOFTWARE: PatentIn Ver
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GENERAL INFORMATION:
APPLICANT: BONAZZI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/245,764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 6374
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4414
LENGTH: 102
                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                             ARLIER FILING DATE: 1998-02-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PPLICANT: EDA, VIDANA
ITTLE OF INVENTION: THREE DIMENSIONAL STRUCTURES AND MODELS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NPPLICANT: Maxwell, Kelly F.
NPPLICANT: Garrett, Thomas P.J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPLICANT: McKenzie,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                             1 EDSGLYQCEAAT 12
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| 234 EDSGLYWCEAAT 245
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nes 11; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/099,994 FILING DATE: 1998-09-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/09245764
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ie, Ian F.C.
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NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
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91.7%;
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Pred. No. 0.041;
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; FEATURE:
; NAME/KEY: SITE
; LOCATION: (349)
; OTHER INFORMATION: X
PCT-US00-05882-1128
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                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 91.7
Matches 11; Conservative
                                                                                                                                            Sequence 3410, Appli GENERAL INFORMATION:
CURRENT APPLICATION NUMBER: US/60/170,373
CURRENT FILING DATE: 1999-12-13
NUMBER OF SEQ ID NOS: 4282
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-702-021-14
                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                              FILE REFERENCE: CL000149
                                                                                                                               APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides FILE REFERENCE: PA106PCT CURRENT APPLICATION NUMBER: PCT/US00/05882 CURRENT FILING DATE: 2000-03-08 EARLIER APPLICATION NUMBER: 60/124,270 EARLIER FILING DATE: 1999-03-12
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Best Local Similarity
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NAME/KEY: SITE
LOCATION: (208)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2000-1
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                                                                                                                                                                                                                                     278
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                                                                                                                            Bonazzi,
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Pred. No.
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0.059;
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; ORGANISM: Human US-60-170-373-3410

TYPE: PRT

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Query Match
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JS-60-177-571-3365
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SEQ ID NO 3365
LENGTH: 81
TYPE: PRT
                            Query Match
Best Local S
Matches 9
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Matches 9; Conserv
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SOFTWARE: FastSEQ for
SEQ ID NO 1676
LENGTH: 93
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CURRENT FILING DATE: 2000-01-27
NUMBER OF SEQ ID NOS: 5082
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CURRENT FILING DATE: 1999-12-28
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                                                                                                                 NAME/KEY: VARIANT
LOCATION: (1)...(93)
OTHER INFORMATION: Xaa = Any Amino Acid
                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: ISOLATED HUMAN ITLE OF INVENTION: ACID MOLECULES ITLE OF INVENTION: THEREOF
                                                                                                                                                                                                                                                                                                                                                                                ENERAL INFORMATION:
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                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                   PPLICANT: BONAZZI, VIVIEN
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                                                                                                                                                                                                                                                                                                        LE REFERENCE:
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                              Local Similarity
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EDSGLYQCEA 10
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90.0%;
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75.08;
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75.08;
                            Score 50; DB Pred. No. 0.25 0; Mismatches
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Pred. No.
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                                                        Length 93;
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Query Match
Best Local Similarity
Thes 9; Conserve
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; ORGANISM: HUMAN
US-60-173-469-1887
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                                                                                                        US-60-173-686-1676
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CCATION: (1)...(93)
OTHER INFORMATION: Xaa =
US-60-173-463-626
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                                                     GENERAL INFORMATION:
APPLICANT: BONAZZI,
                                                                                     Sequence 1676, Application US/60173686
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1887
LENGTH: 93
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SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 626
LENGTH: 93
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/173,469
CURRENT FILING DATE: 1999-12-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTIENS AND
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT FILING DATE: 1999-12-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN ION CHANNEL PROTEINS,
TITLE OF INVENTION: WUCLEIC ACID MOLECULES ENCODING HUMA
TITLE OF INVENTION: AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: BONAZZI,
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                                                                                                                                                                                                                                                      Conservative
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               ISOLATED HUMAN KINASE PROTEINS, NUCLEIC ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES
                                                     VIVIEN
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Pred. No. 0
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APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN KINASE PI
TITLE OF INVENTION: ACID MOLECULES ENCODING
TITLE OF INVENTION: THEREOF
FILE REFERENCE: CLO00182
CURRENT APPLICATION. UNMER: US/09/474,434
CURRENT FILING DATE: 1999-12-28
NUMBER OF SEQ ID NOS: 1872
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1311
                                                                                                                                                            RESULT 13
US-09-474-434-1311
Sequence 1311, Application US/09474434; GENERAL INFORMATION:
GENERAL TOWNS PONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               : LOCATION: (1)...(93)
: OTHER INFORMATION: Xaa = Any Amino
US-60-186-282-885
                                                                                                                                                                                                                                                                                                                                                                                           Query Match 78.
Best Local Similarity 90.
Matches 9; Conservative
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SOFTWARE: FRATSEQ for Windows Version 4.0
SEQ ID NO 885
LENGTH: 93
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Sequence 885, Application US/60186282.
GENERAL INFORMATION:
APPLICANT: BODAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS, AND
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: CL000182
CURRENT APPLICATION NUMBER: US/60/173,686
CURRENT FILING DATE: 1999-12-29
NUMBER OF SEQ ID NOS: 1872
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: VARIANT LOCATION: (1)....(93) OTHER INFORMATION: X -60-173-686-1676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/186,282
CURRENT FILING DATE: 2000-03-01
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90.0%;
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Pred. No. 0.25;
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Pred. No. 0:25;
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B 8

70 EDSGTYQCEA 79

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Query Match
Best Local Similarity
---hes 9; Conserv
                                                                                                     ; ORGANISM: HUMAN US-60-173-686-1311
                                                                                                                                     NUMBER OF SEQ ID NOS:
SOFTWARE: FASTSEQ for
SEQ ID NO 1311
LENGTH: 95
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                   US-60-173-686-1311
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT
; ORGANISM: HUMAN
US-09-474-434-1311
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Best Local Similarity you
Thes 9; Conservative
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                                                                                                                                                                                                                                     TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTIENS AND USES FILE REFERENCE: CL000182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1481, Application US/60173469
                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/173,686
CURRENT FILING DATE: 1999-12-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/60/173,469
CURRENT FILING DATE: 1999-12-29
NUMBER OF SEQ ID NOS: 2120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: BONAZZI, VIVIEN
TITLE OF INVENTION: ISOLATED HUMI
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 95
TYPE: PRT
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1 EDSGLYQCEA 10
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                                          78.1%;
90.0%;
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90.0%;
                                            Score 50;
Pred. No.
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Pred. No. 0.25;
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Pred. No. 0.25
0; Mismatches
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                                                       DB 23; Length 95;
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Search completed: June 4, 2001, 12:20:04 Job time: 517 sec

GenCore version Copyright (c) 1993 - 2000

4.5 Compugen Ltd.

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Minimum
Maximum
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Maximum Match 100%
Listing first 45 summaries
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5 US-09-270-849B-192970
5 US-09-270-849B-183931
1 PPCT-US01-09226-47
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PCT-US01-11988-2222
US-09-833-245-2222
US-09-814-950-4
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US-09-809-991-759
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Sequence 10, Appl
Sequence 6, Appl
Sequence 760, App
Sequence 759, App
Sequence 759, App
Sequence 188150,
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Sequence 3, Appli	Sequence 19, Appl	Sequence 17, Appl	Sequence 15, Appl	Sequence 8, Appli	Sequence 5, Appli.	Sequence 6, Appli	Sequence 20, Appl	Sequence 14, Appl	Sequence 13, Appl	Sequence 186367,	Sequence 186137,	Sequence 182954;	Sequence 190337,	Sequence 5, Appli	Sequence 3, Appli	Sequence 14, Appl	Sequence 7, Appli	

## ALIGNMENTS

RESULT 1 US-09-284-107-26

Sequence 26, Application US, GENERAL INFORMATION:
APPLICANT: Logtenberg, Ton APPLICANT: de Kruif, Corn

Application US/09284107

APPLICANT: de Kruif, Cornelis Adriaan John TITLE OF INVENTION: METHODS AND MEANS FOR SELECTING PEPTIDES TITLE OF INVENTION: AND PROTEINS HAVING SPECIFIC AFFINITY FOR A TARGET FILE REFERENCE: 313632000600

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RESULT 2
PCT-US01-10462-11
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                          Sequence 11, Application GENERAL INFORMATION:
                                                                           APPLICANT:
APPLICANT:
APPLICANT:
                            APPLICANT: Liu, Chenghua
APPLICANT: Tang, Y. Tom
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Radoje T.
TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR B
TITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
                                                                                                                          APPLICANT: HYSEQ,
APPLICANT: Yamaza
URRENT APPLICATION NUMBER: PCT/US01/10462
                                                                                                                                                                                                                                                       1 EDSGLYQCEAAT 12
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1 EDSGLYQCEAAT 12
                                                                                                           Yamazaki, Vici
Asundi, Vinod
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; OTHER INFORMATION: Oligopeptides representing the extracellular; OTHER INFORMATION: domain of CD64 US-09-284-107-26

NUMBER OF SEQ ID NOS: 29

SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 26

LENGTH: 12

LENGTH: 12

ORGANISM: Artificial Sequence FEATURE:

CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08

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Query Match
Best Local Similarity
8; Conserve
                                                                                                                           CT-US01-10462-6
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PCT-US01-10462-11
                                                                                       GENERAL INFORMATION:
                                                                                                      Sequence 6, Application PC/TUS0110462
                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 10
LENGTH: 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 14
SOFTMARE: PatentIn version 3.0
SEQ ID NO 11
LENGTH: 103
                                    APPLICANT:
                                                                      APPLICANT: HYSEQ, INC.
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 09/784,748
PRIOR FILING DATE: 2001-02-14
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 09/540,217
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEC. T.
                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     equence 10, Application PC/TUS0110462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/784,748
PRIOR FILING DATE: 2001-02-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      URRENT APPLICATION NUMBER: PCT/US01/10462
URRENT FILING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PPLICANT: Tang, Y. Tom
PPLICANT: Drmanac, Radoje T.
ITLE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR BING
ITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
ILE REFERENCE: 21272-051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
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                                                                                                                                                                                                                                                                          Local Similarity es 8; Conserv
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                                                                                                                                                                                                                     EDSGLYQCEAA 11
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                                Yamazaki, Victoria
Asundi, Vinod
                   Liu, Chenghua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Asundi, Vinod
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                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                         73.4%;
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Pred. No. 0.46
1; Mismatches
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Pred. No.
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                                                                                                                                                                                                                                                                                                        DB 1;
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US-09-809-391-759
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                                                                                                                        GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: 186 Human Secreted proteins
FILE REFERENCE: PZ002P2
SEQ ID NO 759
LENGTH: 198
                                                                                                                                                                                                                                                                                                                                                         Sequence 759,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 760
                                   Prior application data removed - NUMBER OF SEQ ID NOS: 761 SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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Best Local
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                                                                                         CURRENT APPLICATION NUMBER: US/09/809,391
CURRENT FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/809;391
CURRENT FILING DATE: 2001-03-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Ruben et al. TITLE OF INVENTION: 186 FILE REFERENCE: PZ002P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prior application data removed NUMBER OF SEQ ID NOS: 761
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TYPE: PRT
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PRIOR FILING DATE: 2000-08-23
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TYPE: PRT
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ITLE OF INVENTION: METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR B

ITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES

ILE REFERENCE: 21272-051
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                                                                                                                                                                                                                                                                                                                                     1 EDSGLYQC 8
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         186 Human Secreted proteins
                                                                                                                                                                                                                                                                                                                                                                                        71.9%; Score 46; 100.0%; Pred. No.
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    consult PALM or file wrapper

                                                                      consult PALM or file wrapper
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0.47;
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; ORGANISM: Homo sapiens US-09-809-391-759

Conservative

0

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Indels

0;

Gaps

71.9%;

DB 5;

Length 198;

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Query Match
Best Local Similarity
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RESULT 9
US-09-142-956B-14
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Best Local Similarity 100.
Conservative
                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 188150
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SEQ ID NO 478
LENGTH: 234
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-270-849B-188150 sequence 188150, Application US/09270849B ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-809-391-478
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                                                                                                                                                                                                                                                                                                                                                                              CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: 186 Human Secreted proteins FILE REFERENCE: PZ002P2
                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                             URRENT APPLICATION NUMBER: US/09/270,849B
                                                                                                                                                                                                                                                                                                                                                                                                                ILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PPLICANT: Ruben et al
                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                               PPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                              ENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                                             TTLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 106 EDSGLYQC 113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                106 EDSGLYQC 113
                                                                                                                                Local Similarity 54.
                                                               2 DSGLYQCEAAT 12
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68 DSGIYECQVST 78
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                                                                                                                                             68.88;
54.58;
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Pred. No. 0.65;
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                                                                                                                                                 Pred. No.
                                                                                                                                Mismatches
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                                                                                                                                                               DB 5;
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                                                                                                                                                               Length 261;
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                                                                                                                                                                                                                            any amino acid
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Sequence 14, Application US/09815626 GENERAL INFORMATION:

APPLICANT: Glucksmann, Maria Alexandra
TITLE OF INVENTION: 33395, A NOVEL HUMAN LEUCINE-RITITLE OF INVENTION: FAMILY MEMBER AND USES THEREOF

HUMAN LEUCINE-RICH REPEAT

CURRENT APPLICATION NUMBER: US/09/815,626
CURRENT FILING DATE: 2001-03-23

URRENT FILING DATE: 2001-03-23 RIOR APPLICATION NUMBER: US 60/191,863

ILE REFERENCE: 10448-031001

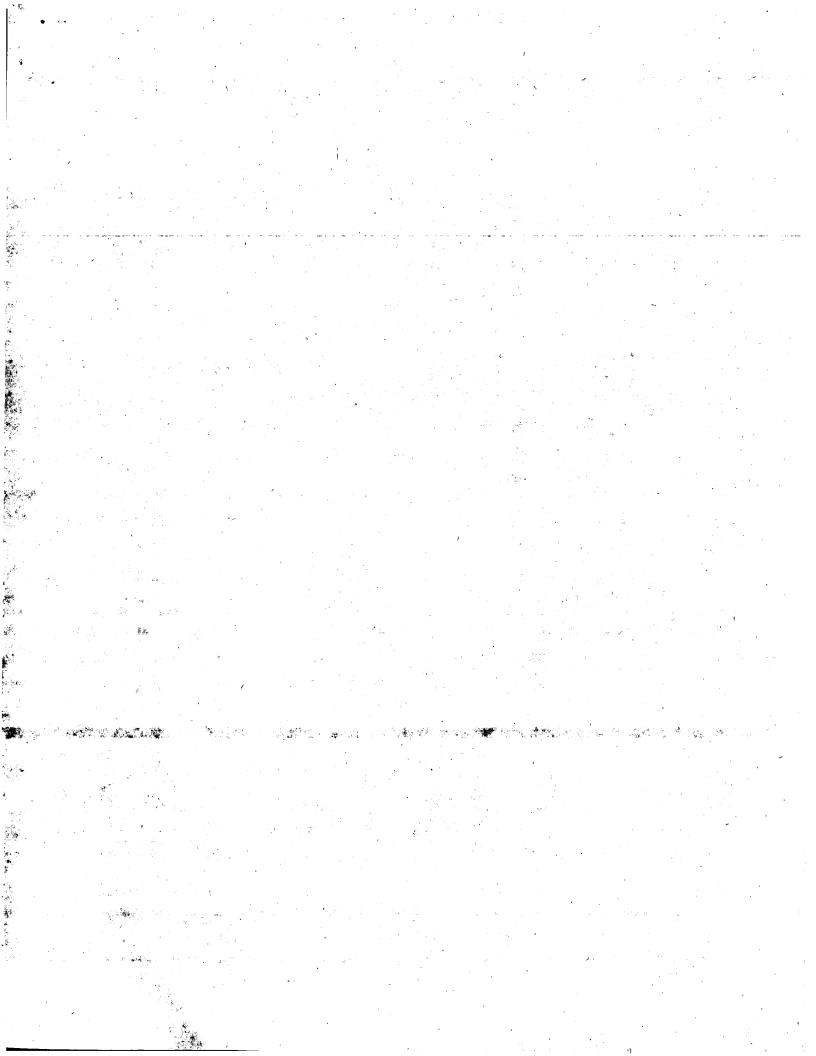
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                                                                                                                   Query Match
Best Local Similarity
7; Conserve
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Best Local Similarity
Thes 7; Conserve
RESULT 11
US-09-815-626-14
                                                                                                                                                                                                                      ; ORGANISM: Homo sapien
US-09-232-773A-15
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JS-09-142-956B-14
                                                                                                                                                                                                                                                                                   SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15, Application US/09232773A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 14, Application US/09142956B
                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/232,773A CURRENT FILING DATE: 1999-07-15
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Kendall, Richard L.
APPLICANT: Thomas, Kenneth A., Jr.
TITLE OF INVENTION: INHIBITOR OF VASCULAR ENDOTHELIAL CELL
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-232-773A-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: JP 9/19706 PRIOR FILING DATE: 1997-01-17
                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE: 1998-01-16
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TITLE OF INVENTION: VEGF-BINDING POLYPEPTIDES
FILE REFERENCE: 06501-021001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                     LENGTH: 789
TYPE: PRT
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                                                               730 EDEGLYTCOACS 741
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                                                                                  1 EDSGLYQCEAAT 12
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                                                                                                                                   Conservative
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                                                                                                                                                     Score 43; [Pred. No. 8;
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Pred. No.
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                                                                                                                                                                   Length 789;
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; OTHER INFORMATION: consensus sequence US-09-822-687-9
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Best Local Similarity
Thes 8; Conserve
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; OTHER INFORMATION: consensus sequence
US-09-815-626-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/822,687
CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: US 60/193,919
PRIOR FILING DATE: 2000-03-31
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO
                                                                                                                                                                                                                                                                                               Sequence 13, Application PC/TUS0110462 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
           CURRENT APPLICATION NUMBER: PCT/US01/10462
CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/784,748
PRIOR FILING DATE: 2001-02-14
PRIOR APPLICATION NUMBER: 09/649,167
PRIOR FILING DATE: 2000-08-23
PRIOR FILING DATE: 2000-08-23
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 9, Application US/09822687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -09-822-687-9
                                                                                                                              APPLICANT: Liu, Chenghua
APPLICANT: Tang, Y. Tom
APPLICANT: Drmanac, Radoje T.
APPLICANT: Drmanac, Radoje T.
APPLICANT: DRMANAC, METHODS AND MATERIALS RELATING TO INSULIN-LIKE GROWTH FACTOR BINITITLE OF INVENTION: PROTEIN-LIKE POLYPEPTIDES AND POLYNUCLEOTIDES
ALL REFERENCE: 21272-051
                                                                                                                                                                                                                                                                                                                                       -US01-10462-13
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APPLICANT: Glucksmann, Maria Alexandra
APPLICANT: THEREOF
                                                                                                                                                                                                                                            PPLICANT:
                                                                                                                                                                                                                                                                             PPLICANT: HYSEQ, INC.
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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
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42 EDSGLYTCVA 51
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 APPLICATION NUMBER:
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Asundi, Vinod
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80.0%;
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80.0%;
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Pred. No. 0.81;
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Qy
; ORGANISM: Homo sapiens
PCT-US01-11988-133
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Best Local Similarity
Thehes 7; Conserv.
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                                              SEQ ID NO 133
LENGTH: 359
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Best Local
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                                                                                            PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
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                                                                                SOFTWARE:
                                                                                                                                           PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                          CURRENT FILING DATE: 2001-01-12
PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: PCT/US01/11988 CURRENT FILING DATE: 2001-01-12
                                                                                                                                                                                                                                             APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 08/657,753
PRIOR FILING DATE: 1996-05-30
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/266,091 CURRENT FILING DATE: 1999-03-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Bennett, C. Frank
TITLE OF INVENTION: ENHANCED EFFICACY OF LIPOSOMAL ANTISENSE THERAPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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Hope, Michael
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Pred. No. 11;
1; Mismatches
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Pred. No.
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	Search completed: June 4, 2001, 12:21:29 Job time: 591 sec		Db 233 DHSGSYWCEAAT 244	Qy 1 EDSGLYQCEAAT 12	Matches 8; Conservative	Query Match 64.1%; Score 41; DB 1; Length 359;
	, 2001,		4		tive	64.18;
*	12:21:29				1; Mismatches	Score 41; DB 1;
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                                                                                                                                             DB
                       3 US-60-160-209-4252
3 US-60-160-203-4414
6 US-09-245-764-7
1 US-09-702-021-14
PCT-US00-05882-1128
US-08-332-562-94
US-08-462-973-94
US-08-462-973-94
                                                                                                                                                                                 SUMMARIES
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Sequence 4252, Ap
Sequence 4414, Appl
Sequence 7, Appl 1
Sequence 114, Appl
Sequence 118, Ap
Sequence 94, Appl
Sequence 94, Appl
Sequence 25, Appl
Sequence 27859, A
Sequence 27859, A
                                                                                                                                         Description
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ALIGNMENTS

ct score:	61 1 EDGNVLKRSPEL 12	
ng table:	BLOSIM62 Gapop 10.0 , Gapext 0.5	
hed:	1009251 seqs, 160854530 residues	,
number of	hits satisfying chosen parameters: 1009251	51
um DB seq um DB seq	length: 0 length: 2000000000	
processing:	: Minimum Match 08 Maximum Match 1008 Listing first 45 summaries	
ase .	Pending_Patents_AA_Main:*  1: /cgn2_6/ptodata/2/paa/PCTUS_COMB.pep:*  2: /cgn2_6/ptodata/2/paa/US06_COMB.pep:*  3: /cgn2_6/ptodata/2/paa/US07_COMB.pep:*  4: /cgn2_6/ptodata/2/paa/US080_COMB.pep:*  5: /cgn2_6/ptodata/2/paa/US080_COMB.pep:*	
	o: /cgn2_6/ptodata/2/paa/US082_COMB.pep:* 7: /cgn2_6/ptodata/2/paa/US084_COMB.pep:* 8: /cgn2_6/ptodata/2/paa/US085_COMB.pep:* 9: /cgn2_6/ptodata/2/paa/US085_COMB.pep:*	
· .	10: /cgn2_6/ptodata/2/paa/USOU86_COMB.pep:* 11: /cgn2_6/ptodata/2/paa/USOU86_COMB.pep:* 12: /cgn2_6/ptodata/2/paa/USOU86_COMB.pep:* 13: /cgn2_6/ptodata/2/paa/USOU86_COMB.pep:*	•
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US-09-489-039A-130 US-09-417-507-4260 US-09-619-049-411 US-60-167-217-7900 US-60-171-627-660

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16492, A 16167, A 16165, Ap 16965, Ap 16965, Ap 13098, A 13098, A 42680, Ap 7900, Ap 660, Ap 660, Ap 6310, Ap 6310, Ap

Sequence Sequence Sequence

7942, 23, A

Sequence Sequence

US-60-173-464-6310
US-60-191-631-7800
US-60-191-637-7800
US-60-191-637-7801
US-60-191-637-36817
US-60-191-637-36817
US-60-167-217-22133
US-60-167-217-22133
US-60-173-464-17987
US-09-733-089-15889
US-09-733-089-15889
US-08-911-503-634
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Sequence 4292, Ap Sequence 634, App Sequence 634, App Sequence 634, App Sequence 137, App Sequence 834, App Sequence 50997, A Sequence 50997, A Sequence 664, App Sequence 664, App Sequence 6322, Ap Sequence 5322, Ap

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Qy 1 EDGNVLKRSDEL 12	Query Match 100.0%; Score 61; DB 23; Length 96; Best Local Similarity 100.0%; Pred. No. 0.00038; Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps	; LOCATION: (1)(96) ; OTHER INFORMATION: Xaa = Any Amino Acid US-60-160-209-4252	; TYPE: PKT ; ORGANISM: HUMAN ; FEATURE: ; NAME/KEY: VARIANT	SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 4252 LENGTH: 96	FILE REFERENCE: CL000113  CURRENT APPLICATION NUMBER: US/60/160,209  CURRENT FILING DATE: 199-10-19  UNMBER OF SEO ID NOS: 4646	NASE PROTEINS, NUCLEIC CODING HUMAN KINASE PROTIENS	RESULT 1 US-60-160-209-4252  G-160-209-4252  G-160-209-4252
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US-09-702-021-14
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; ORGANISM: Homo sapiens
US-09-245-764-7
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US-60-160-203-4414
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                                                         GENERAL INFORMATION:
                                                                     Sequence 14, Application US/09702021
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Best Local Similarity
Matches, 12; Conserv
APPLICANT: C. Fraser
TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF
FILE REFERENCE: 7853-217
                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 6374
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4414
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GENERAL INFORMATION:
APPLICANT: BONAZZI, VIVIEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/60/160,203
CURRENT FILING DATE: 1999-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: ISOLATED HUMAN SECRETED PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN SECRETED PROTEINS AND
FILE REFERENCE: CL000116
THERENT APPLICATION
                                                                                                                                                                                                                                                                                                                                                                                           OFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                           ARLIER APPLICATION NUMBER: 60/073,972
ARLIER FILING DATE: 1998-02-06
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TYPE: PRT
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                                                                                                                                                         246 EDGNVLKRSPEL 257
                                                                                                                                                                                                                           Local Similarity nes 12; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87 EDGNVLKRSPEL 98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/09/245,764 FILING DATE: 1999-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7, Application US/09245764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                             LING DATE: 1998-09-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Powell, Maree S.
McKenzie, Ian F.C.
                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CATION NUMBER:
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Pred. No.
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Pred. No.
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Query Match
Best Local Similarity
""+"hes 12; Conserv
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                                                                                                                                                                                                                                                                                                                          Best
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SEQ ID NO 1128
LENGTH: 399
                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ fo:
SEQ ID NO 14
LENGTH: 374
                                                                                                                                                                                            -08-332-562-94
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT FILING DATE: 2000-03-08 EARLIER APPLICATION NUMBER: 60/EARLIER FILING DATE: 1999-03-12
                                                                                                                                                              SENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: PCT/US00/05882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/702,021 CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Human Cancer Associated FILE REFERENCE: PA106PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                           FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: SITE
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                                                                                                                                                APPLICANT:
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                            CORRESPONDENCE ADDRESS
                                                                                                    APPLICANT:
                                            UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                   290 EDGNVLKRSPEL
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                                                                                                                                                                                                                                                                                                           Local Similarity
mes 12; Conser
                                                                                                                                                                                                                                                                              1 EDGNVLKRSPEL 12
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                                                           INVENTION:
                                                                                                                                                                           Application US/08332562
: Foley & Lardner 3000 K Street, N.W.,
                                                                                   BONADONNA, Lisa
                                                                                                  SCOTT, Bernadette M
                                                                                                                HIBBS, Margaret L.
                                                                                                                                             MCKENZIE,
                                                      HULETT, Mark D.

ENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                 HOGARTH, Mark
                                                                                                                                                                                                                                                                                                           Conservative
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                    Length 399;
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STREET:

Suite 500

COUNTRY:

20007-5109

CITY: Washington STATE: D.C.

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Best Local Similarity
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SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                quence 94, Applicat
ENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
 FILING DATE: 05-JUN-1335
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/332,562
                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                         ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                        UMBER OF SEQUENCES:
                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                      CITY: Washington STATE: D.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                                     STREET:
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FILING DATE: 31-OCT-1994
                                                                                      APPLICATION NUMBER: US/08/462,973
                                                                                                                                                                                                                        COUNTRY:
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                                                                                                                                                                                                                                                                                       ADDRESSEE:
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                                                                                                                                                                                                        20007-5109
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                                                                                                                                                                                                                                                                     3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                                                                                                                                         HOGARTH, Mark P.
HIBBS, Margaret L.
                                                                                                                                                                                                                                                                                                                                                                                                                                           MCKENZIE,
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                                                                                                                                                                                                                                                                                                                                        HULETT, Mark D.

TENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                  PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                           BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                         Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stephen A.
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31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                             Bernadette M.
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83.3%;
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Pred. No.
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Best Local Similarity
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                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU90/00513
FILING DATE: 25-OCT-1990
                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
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REFERENCE/DOCKET NUMBER: 16.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEPAX: (202)672-5399
TELEPAX: (202)672-5399
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 'AU PJ 7045
FILING DATE: 25-OCT-1989
ATTORNEY/AGENT INFORMATION:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: HYBRID FC RECEPTOR MOLECULES NUMBER OF SEQUENCES: 29
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                                                                                                                       FILING DATE: 19 CLASSIFICATION:
                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 EDGNVLKRSPEL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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26-MAY-1986
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27-MAY-1992
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Pred. No. 0
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                                                                                                                                                                                         Version #1.25
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Query Match
Best Local Similarity
6; Conserv
; ORGANISM: DROSOPHILA
US-60-191-681-22468
                                                                                                                                                                                                                                                                                              RESULT 10
US-60-191-681-22468
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JS-60-191-637-27859
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                                                  NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ fo:
SEQ ID NO 22468
LENGTH: 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 27859
LENGTH: 99
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Best Local Similarity
Matches 10; Conserv
                                                                                                                            CURRENT APPLICATION NUMBER: US/60/191,681
CURRENT FILING DATE: 2000-03-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -60-191-637-27859
                                                                                                                                                                               TITLE OF INVENTION:
FITLE OF INVENTION:
FITLE OF INVENTION:
                                                                                                                                                                                                                                                             equence 22468, Application US/60191681 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/60/191,637
                                                                                                                                                                     TLE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VERLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM TITLE OF INVENTION: GENE SEQUENCES EXPRESSED FROM THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION: HERE THE OF INVENTION:
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TYPE: AMINO ACID
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                                                                                                                                                                                                                                         Li, Peter, W.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                         for Windows Version 4.0
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                                                                                                                                                                                     USES THEREOF.
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54.5%;
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83.3%;
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Pred. No.
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0.36;
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Query Match

68.9%;

Score 42;

DВ

23;

Length 99;

US-09-248-796-16492

Sequence 16492, Application US/09248796 GENERAL INFORMATION:

APPLICANT: Keith Weinstock et al TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBI

RESULT 13

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                                                                                                                                            ; LENGTH: 407
TYPE: PRI
; ORGANISM: Arabidopsis thaliana
US-09-347-801-23
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US-60-167-217-7942
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                                                              Matches
                                                                                           Query Match
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7942
LENGTH: 299
TYPE: PRT
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APPLICANT: L1, Petcar W. D.
TITLE OF INVENTION: ISOLATED DROSOPHILA PROTEINS,
TITLE OF INVENTION: ACID MOLEUCLES ENCODING DROSC
TITLE OF INVENTION: THÉREOF
FILE REFERENCE: CL000152
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Best Local
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Matches 6; Conserv
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                                                                                                                                                                                                                                        EARLIER APPLICATION NUMBER: 60/092,438 EARLIER FILING DATE: July 10, 1998 NUMBER OF SEQ ID NOS: 23
                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/347,801 CURRENT FILING DATE: 1999-07-02
                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Plant Protein Kinases FILE REFERENCE: BB-1171
                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Allen, Steve APPLICANT: Lee, Jian Mi
                                                                                                                                                                                                                          SOFTWARE: Microsoft Office 97
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/60/167,217
CURRENT FILING DATE: 1999-11-24
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21
                            2 DGNVLKRSPEL 12
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DGDALKRRPEL 31
                                                             Similarity 72.
8; Conservative
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                                                                            68.9%;
72.7%;
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54.58;
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                                                                        Score 42;
Pred. No.
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Pred. No.
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                                                                          DB 17;
13;
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                                                                                        Length 407;
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NUMBER OF SEQ ID NOS: 7544
SEQ ID NO 7543
LENGTH: 166
TYPE: PRT
ORGANISM: S.epidermidis
US-09-450-969-7543
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US-09-248-796-16167
US-09-248-796-16167, Application US/09248796
; Sequence 16167, Application US/09248796
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; APPLICANT: APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICANT APPLICAN
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Best Local Similarity
Thes 7; Conserve
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US-09-248-796-16167
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Best Local Similarity
Matches 6; Conserv
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GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: PATH99-09A
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SEQ ID NO 16167
LENGTH: 490
TYPE: PRT
                                                                                                                                               Matches
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Best Local Similarity
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CURRENT FILING DATE: 1999-11-29
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CURRENT FILING DATE: 1999-02-12
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EQ ID NO 16492
LENGTH: 368
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330 KNGNIFKLSPEI 341
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1:||::||
109 EEGNIIKRKTGVLMLNPEL 127
                                                                       1 EDGNVLKR----,---SPEL 12
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                                                                                                                                               Conservative
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58.3%;
                                                                                                                                       Score 37.5; D
Pred. No. 32;
4; Mismatches
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                       DB 18;
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Search completed: June 4, 2001, 12:20:04 Job time: 517 sec

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Result
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Maximum DB
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2000
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/cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep:*
/cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep:*
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US-09-270-849B-192159
US-09-270-849B-191599
US-09-549-066-37
US-09-549-066-36
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PCT-US01-04098A-3250
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PCT-US01-04098A-1283
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US-09-739-449-
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US-09-549-066-2
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US-09-739-449-11789
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PCT-US01-04098A-3229
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                    US-09-549-066-35
US-09-327-563B-4
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•	Sequence 874, App	Sequence 529, App	Sequence 1162, Ap	Sequence 723, App	Sequence 1, Appli	Sequence 2, Appli	Sequence 13, Appl	Sequence 3, Appli	Sequence 1027, Ap	Sequence 14, Appl	Sequence 6, Appli	Sequence 10210, A	Sequence 8764, Ap	Sequence 595, App	Sequence 185071,	Sequence 85, Appl	Sequence 1259, Ap	Sequence 2, Appli

### ALIGNMENTS

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US-09-739-449-11283
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US-09-284-107-27
                                                                                                                                                                                                                        Sequence 11283, Appl GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ'ID NO 27
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 27, Application US/09284107 GENERAL INFORMATION:
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 11283
                                          CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR ETLING DATE: 2000-02-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/284,107
CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
                                                                                                                                 APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens
FILE REFERENCE: 38-10(15490)C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: EP 96202791.8 PRIOR FILING DATE: 1996-10-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: de Kruif, Cornelis Adriaan John
TITLE OF INVENTION: METHODS AND MEANS FOR S
TITLE OF INVENTION: AND PROTEINS HAVING SP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Logtenberg, Ton APPLICANT: de Kruif, Corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 313632000600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE: Oligopeptides representing the extracellular OTHER INFORMATION: Oligopeptides representing the extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                     1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                                                                                                                                                1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                             Application US/09739449
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Pred. No.
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2e-05;
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LENGTH: 124

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1 EDGNVLKRSPEL 12

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Query Match
Best Local Similarity
7; Conserve
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Best Local Similarity
7; Conserv
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US-09-270-849B-183641
; Sequence 183641, Application US/09270849B
                                                                                                  ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: at any location having Xaa, Xaa means any amino acid
US-09-270-849B-183641
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Best Local Similarity
Tatches 7; Conserve
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                                                                                                                                                   SEQ ID NO 133
LENGTH: 359
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LENGTH: 189
                                                                                                                                    TYPE: PRT
                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING.DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: PCT/US01/11988 CURRENT FILING DATE: 2001-01-12 PRIOR APPLICATION NUMBER: 60/229, 358 PRIOR PILING DATE: 2000-04-12 PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR APPLICATION NUMBER: 60/256, 931 PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
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TITLE OF INVENTION: Insect genome survey devices
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/270,849B
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                                                                                                                                                                                                                                                                                                                                                                                                   ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                 PPLICANT: Human Genome Sciences, Inc.
ITLE OF INVENTION: Albumin Fusion Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                       LE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             108
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                               59.0%;
58.3%;
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          Score 36; DB Pred. No. 73; 2; Mismatches
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Pred. No.
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                                               1;
                                            Length 359;
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Вþ
                                                                                                                               FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic US-09-270-849B-192159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QΥ
                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-270-849B-192159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
US-09-833-245-133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-833-245-133
                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 192159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 58.
Matches 7; Conservative
                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 192159, Application US/09270849B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver.
SEQ ID NO 133
LENGTH: 359
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 133,
                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Swimmer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/199, 384
PRIOR FILING DATE: 2000-04-25
NUMBER OF SEQ ID NOS: 2267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/833,245
CURRENT FILING DATE: 2001-04-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Human Genome Sciences, Inc: TITLE OF INVENTION: Albumin Fusion Proteins
                                                                                                                                                                                                    LENGTH: 193
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/229, 358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: PF546PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      245 EDNQVWKQSPQL 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 245 EDNQVWKQSPQL 256
                1 EDGNVLKRSP 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 EDGNVLKRSPEL 12
EDGVVLRRPP 15
                                                                 h 57.4%;
Similarity 70.0%;
7; Conservative
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                                                                                                                                                                                                                                                                                                                                                             Insect genome survey devices
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          59.0%;
58.3%;
                                                               Score 35; DB
Pred. No. 54;
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                DB 5;
54;
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                                                                                              Length 193;
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                                                                 Indels:
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US-09-270-849B-191599
; Sequence 191599, Application US/09270849B
; GENERAL IMPORMATION:
; APPLICANT: Swimmer et al.
; TITLE OF INVENTION: Insect genome survey devices

SEQ ID NO 191599

LENGTH: 470

TYPE: PRT

FEATURE:

ORGANISM: Artificial Sequence

Description of Artificial Sequence: Synthetic

NUMBER OF SEQ ID NOS: 1 SOFTWARE: PatentIn Ver.

CURRENT APPLICATION NUMBER: US/09/270,849B CURRENT FILING DATE: 1999-03-17

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Query Match
Best Local Similarity
...thes 6; Conserv
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US-09-549-066-37
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                                                                                                         TOPOLOGY:
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OTHER INFORMATION:
US-09-270-849B-191599
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Best Local Similarity
                                                                                                                                                                                NFORMATION FOR SEQ ID NO:
                                                                                                                                                  SEQUENCE CHARACTERISTICS
LENGTH: 497 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Re
                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 13-APRI
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                        NTTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IIILE OF INVENTION: Programmed Cell Death Genes and Proteins NUMBER OF SEQUENCES: 85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        282 EDGVVLRRPP 291
                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 35
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: FILING DATE: 10-JU
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington
                                                                                                                                       TYPE:
                                                                                                                                                                                                                               TELEPHONE:
                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/080,850 FILING DATE: 24-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/549,066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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1 EDGNVLKRSPEL 12
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                                                                                                                                                                                                  248636 SSK
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                                                                                                                                                    497 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yuan, Junying
                                                                                                                       both
                                                                                                                                                                                                                                                                                                                                                                                                                                              PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                               (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                   13-APRIL-2000
                                                                                                                                                                                                                                                                                                                                                      10-JUN-1994
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                                            57.4%;
50.0%;
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70.0%;
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                                                                                                                                                                                                                                                             0609.3920001
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Pred. No. 1.6e+02;
                                              Score 35;
Pred. No.
                               Mismatches
                                            7e+02;
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Query Match
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                                                                                                                                                                                                                                                          RESULT 10
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                                                   SOFTWARE: PatentIn Ver. SEQ ID NO 1066
                                                                                                                                                                                                                        Sequence 1066, Application PC/TUS0101312
                                                                                    Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510
                                                                                                                      CURRENT APPLICATION NUMBER: PCT/US01/01312
CURRENT FILING DATE: 2001-05-09
                                                                                                                                                                      APPLICANT: Human Genome Sciences, Inc., et al. TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
                                                                                                                                                         FILE REFERENCE: PTZ32PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NFORMATION FOR SEQ
ORGANISM: Homo sapiens
                                   ENGTH:
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APPLICATION NUMBER: US 08/080,850
FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMATION:
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465 QGANILKQMPEL 476
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SOFTWARE: Patenti
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FILING DATE: 13-APRIL-2000
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                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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FEATURE: NAME/KEY: SITE

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; ORGANISM: Agrobacterium tumefaciens US-09-739-449-12341
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; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
PCT-US01-01312-1066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 7; Conser
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Best Local Similarity 60.0
Watches 6; Conservative
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SEQ ID NO 12341
LENGTH: 216
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APPLICANT: Slater, Steven C.
TITLE OF INVENTION: AGIODACTERIUM 'tumefaciens Genome
FILE REFERENCE: 38-10(15490)C
CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
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                                                                                                                                   RRIOR APPLICATION NUMBER: 09/654,936-
PRIOR FILING DATE: 2000-09-01
PRIOR APPLICATION NUMBER: 09/620,325
RIOR FILING DATE: 2000-07-19
                                                                                                                                                                                                                                                                     JURRENT APPLICATION NUMBER: PCT/US01/04098A
JURRENT FILING DATE: 2001-02-05
PRIOR APPLICATION NUMBER: Not Yet Assigned
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 09/728,422
PRIOR APPLICATION NUMBER: 09/728,422
PRIOR APPLICATION NUMBER: 09/693,325
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                                                                                              RIOR FILING DATE: 2000-06-20
                                                                                                                                                                                                               PRIOR FILING DATE: 2000-10-20
RIOR APPLICATION NUMBER: 09/663,561
PRIOR FILING DATE: 2000-09-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   equence 1261, Application PC/TUS0104098A
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                                                                                                                                                                                                                                                                                                                                                                                                                        ILE REFERENCE: 21272-029
                                                                                                                                                                                                                                                                                                                                                                                                                                   ITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        180 GNWLKRSP 187
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              APPLICATION NUMBER: 09/560,875
FILING DATE: 2000-04-27
APPLICATION NUMBER: 09/496,914
FILING DATE: 2000-02-03
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QF
SEQ'ID NOS:
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87.5%;
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60.0%;
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Pred. No.
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Pred. No.
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PCT-US01-11797-29

Sequence 29, Application PC/TUS0111797 GENERAL INFORMATION:

APPLICANT: SMITHKLINE BEECHAM CORPORATION APPLICANT: SMITHKLINE BEECHAM p.l.c. TITLE OF INVENTION: NOVEL COMPOUNDS FILE REFERENCE: GP50022 CURRENT APPLICATION NUMBER: PCT/US01/11797 CURRENT FILING DATE: 2001-04-11

PRIOR APPLICATION NUMBER: 60/196,603

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                                   Qy
                                                                                                                                                         ; ORGANISM: Homo sapiens
PCT-US01-04098A-3229
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PCT-US01-04098A-3229
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Best Local Similarity 4/...
"""+" 9; Conservative
                                                                       Matches
                                                                                                    Query Match
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SEQ ID NO 3229
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SEQ ID NO 1261
                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 3960
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides FILE REFERENCE: 21272-029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: Not Yet Assigned PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                           LENGTH: 232
118 EDGSVLKQFLSETEKMSPE 136
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                                                                     Local Similarity hes 9; Conserva
                                   1 EDGNVLKR-----SPE 11
                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 2000-09-01
APPLICATION NUMBER: 09/620,325
FILING DATE: 2000-07-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2000-11-30
APPLICATION NUMBER: 09/693,325
                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 09/560,875
FILING DATE: 2000-04-27
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                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 09/496,914
                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 09/598,075
                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 09/654,936
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 09/663,561 FILING DATE: 2000-09-15
                                                                                                                                                                                                                                                                                                                                              FILING DATE: 2000-06-20
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                                                                     Conservative
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                                                                                  55.7%;
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47.4%;
                                                                                    Score 34; DB 1;
Pred. No. 1e+02;
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Pred. No.
                                                                     Mismatches
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                                                                                                    Length 232;
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RESULT 15
US-09-739-449-11837
Sequence 11837, Application US/09739449
GENERAL INFORMATION:
APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences and Uses Thereof
FILE REFERENCE: 38-10(15490)C
CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 11837
LENGTH: 184
TYPE: PRI
CORGANISM: Agrobacterium tumefaciens
US-09-739-449-11837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; PRIOR FILING DATE: 2000-04-13
; PRIOR APPLICATION NUMBER: 60/199,417
; PRIOR FILING DATE: 2000-04-24
; NUMBER OF SEQ ID NOS: 48
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 29
; LENGTH: 801
; TYPE: PRT
ORGANISM: Homo sapiens
PCT-US01-11797-29
Search completed: June 4, 2001, 12:21:29
Job time: 591 sec
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Best Local Similarity 66.7
Matches 6; Conservative
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Best Local Similarity 54.5%;
Matches 6; Conservative
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133 DGLILKRGP 141
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result No.	Score	Query Match	Query Match Length DB	BDB	ID .	Description
<b>—</b>	60	100.0	374	21	US-09-702-021-14	Sequence 14, Appl
N	60	100.0	399	Ъ	PCT-US00-05882-1128	Sequence 1128, Ap
ω	43	71.7	165	1	PCT-US01-03080-9	Sequence 9, Appli
4	43	71.7	195	┙	PCT-US01-03080-8	Sequence 8, Appli
տ	39	65.0	214	23	US-60-258-251-43	Sequence 43, Appl
6	39	65.0	338	18	8 US-09-467-008-6	Sequence 6, Appli
7	39	65.0	380	ω	US-07-922-466-6	Sequence 6, Appli
œ	39	65.0	380	ω	US-07-922-466A-6	Sequence 6, Appli
9	39	65.0	380	7	US-08-394-442-6	Sequence 6, Appli
10	39	65.0	380	œ	US-08-416-478-6	Sequence 6, Appli

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# ALIGNMENTS

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                             PCT-US00-05882-1128
: Sequence 1128, Application PC/TUS0005882
: GENERAL INFORMATION:
                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens
US-09-702-021-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1
US-09-702-021-14
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SEQ ID NO 14
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Best Local Similarity
Matches 12; . Conserv
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APPLICANT: Craig Rosen, APPLICANT: Steve Ruben
                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: C. Fraser TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE:
                                                                                                                                                                                                                                                                                                 LENGTH: 374
TYPE: PRT
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    Conservative

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RESULT 4
PCT-US01-03080-8
; Sequence 8, App
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                                                                                                                                                                                                                            PCT-US01-03080-9
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LENGTH: 399
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Best Local Similarity 100.0%;
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: PCT/US01/03080 CURRENT FILING DATE: 2001-01-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Human Cancer Associated Gene Sequences and Polypeptides
                                                                                                                                                                                                                                      NAME/KEY: SITE LOCATION: (158)
OTHER INFORMATION: Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: \15
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                       NAME/KEY: SITE
LOCATION: (126)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
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Application PC/TUS0103080
                                                                                                                                                                                                                                                                                                                                                                                                                   PatentIn
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"ENTION: Bcl-2-like Polynucleotides, Polypeptides, and Antibodies
                                                                                                                                           Conservative
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; OTHER INFORMATION: Xaa = Any Amino Acid US-60-258-251-43
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RESULT 6
US-09-467-008-6
; Sequence 6, Application US/09467008
; GENERAL INFORMATION:
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                                                                                                                                                      Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                      SEQ ID NO 43
LENGTH: 214
TYPE: PRT
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/60/258,251
CURRENT FILING DATE: 2000-12-27
NUMBER OF SEQ ID NOS: 102
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 2000-0
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: PCT/US01/03080 CURRENT FILING DATE: 2001-01-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Beasley,
                                                                                                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: CL0001039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                            FEATURE: NAME/KEY: VARIANT LOCATION: (1)...(
                                                                                                                                                                                                                                                                                                         ORGANISM: HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PPLICANT: Human Genome Sciences, Inc. ITLE OF INVENTION: Bcl-2-like Polynucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ILE REFERENCE:
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                                                                                             48
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                                                                                             ELRVLGKDLPRP 59
                                                                                                                          ELQVLGLQLPTP 12
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8; Conser
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66.7%;
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66.7%;
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                                                                                                                                                                       Score 39;
Pred. Nó.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 43;
Pred. No.
                                                                                                                                                         Mismatches
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24;
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CORRESPONDENCE ADDRESS ROMAGNANI, Serg APPLICANT: TRIEBEL, Frederic

SEQUENCES:

MASTRANGEL1,

Sergio Renato

SPLICE VARIANTS

ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C. STREET: 624 \ Ninth Street, N.W., Suite

ZIP: 20001 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COUNTRY: USA

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UKGANISM: Homo sapiens;
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-09-467;008-6
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Best Local Similarity
Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                quence 6, Application US/07922466 ENERAL INFORMATION:
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                                                    SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/922,466
                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                                                                                                                                                                                                                      UMBER OF SEQUENCES:
CLASSIFICATION: 424
TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                            STREET:
                                                                                                                                                                                                      COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: EP 97401404.5
FILING DATE: 18-JUN-1997
APPLICATION NUMBER: PCT/EP98/03307
FILING DATE: 03-JUN-1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/467,008 FILING DATE: 20-Dec-1999 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                       20036
                                                                                                                                                                                                                         Washington : D.C.
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                                                                                                                                                                                                                                                              E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Avenue
                                                                                                                                                                                                      U.S.A.
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                                      19920710
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Frederic
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72.7%;
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257 LTVLGLEPPTP 267

2 LQVLGLQLPTP 12

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                                                                                   US-07-922-466A-6
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US-07-922-466A-6
 Matches
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               Query Match
Best Local Similarity
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 6:
                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 380 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (202)466-0800
                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 07
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)371-2600
                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge A
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TITLE OF INVENTION:
                                                                                               TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                   TYPE:
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STREET: 1100 New York Avenue, Suite 600
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                                                                                                                                                 LENGTH:
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                                                                                                                                               380 amino acids
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 Conservative
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                                                                                                                                                                                                 (202) 371-2540
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72.7%;
                 65.0%;
72.7%;
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Lymphocytes, DNA Sequences Encoding These Proteins And
Lymphocytes, DNA Sequences Encoding These Proteins And
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1; Mismatches
                 Score 39; DB 3;
Pred. No. 2.2e+02;
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Pred. No. 2.2e+02;
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                                Length 380;
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Gaps
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JS-08-394-442-6
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein -08-394-442-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (202) 628-5197
TELEFAX: (202) 737-3528
[INFORMATION FOR SEQ ID NO: 6:
COMPUTER: IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                        257 LTVLGLEPPTP 267
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                                                                                                                                                                   JMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: BROWDY, ROGER L. REGISTRATION NUMBER: 25, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
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                                                                COUNTRY:
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                                                                                   D.C.
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                                                                                                                 1100 New York Avenue,
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                                                                  U.S.A.
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                                                                                                                                 Sterne, Kessler, Goldstein & Fox
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lymphocytes, DNA Sequences Encoding These Proteins And Their Pharmaceutical And Biological Uses
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                                                                                                                                                                             Lymphocytes, DNA Sequences Encoding These Proteins And Their Pharmaceutical And Biological Uses
                                                                                                                                                                                                                                                                                                                                                                                                                                     65.0%;
                                                                                                                                                                                                                New Proteins Produced By Human
                                                                                                                                                                                                                                               Thierry
                                                                                                                                                                                                                               Frederic
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 39; DB 7;
Pred: No. 2.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                    Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 380;
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Query Match
Best Local Similarity
"Atches 8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (202) 833-87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
FILING DATE: 08-SEP-
                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
REFERENCE/DOCKET NUMBER: HERCEND=1B TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 628-5197
                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                   PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                257 LTVLGLEPPTP 267
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Lewis, James L. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
                                                  REGISTRATION NUMBER: 37,971
                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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                                                                                                                                                    CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                  20004
                                                                                                                                                                                                                                                                                                                                                                                                                Washington
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amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   BROWDY AND NEIMARK, P.L.L.C.
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                                                                                                                                                                    04-APR-1995
                                                                                                                                                                                                                                                                                                                                 Floppy disk /
                                                                                                                                                                                                                              UMBER: US/08/474,988
07-JUN-1995
                                                                                                    JMBER: US 07/854,644
08-SEP-1992
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72.7%;
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LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS AND
THEIR PHARMACEUTICAL AND BIOLOGICAL USES
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US-08-474-988A-6
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Best Local Similarity
Matches 8; Conserv
                                         Query Match
Best Local Similarity
                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             equence 6, Application US/08474988A GENERAL INFORMATION:
                                                                                                                                                                                                       NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (202) 737-3528 NFORMATION FOR SEQ ID NO: 6:
                                                                                                                             TYPE: amino acids
TYPOLOGY: 1:--
YOLDGY: 1:--
YOLDGY: 1:--
                                                                                                                                                                                                                                     REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HE
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/854,644
PILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/416,478
FILING DATE: 04-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street, N.W., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 419 Seven
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2 LQVLGLQLPTP 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         : 380 amino acids amino acid
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                                                                                                                                                                                                                                                                                                        YUN, Allen C.
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                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Pred: No. 2.2e+02;
                                              Score 39;
Pred. No. 2
                               Mismatches
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                                             DB 8;
2.2e+02;
                                                            Length 380;
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                               Indels
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UKGANISM: Homo sapiens;
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-467-008-4
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                                                                                                                                   US-08-394-442-7
                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                             Sequence 7, Application:
                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 4:
                   TITLE OF INVENTION:
                                                         APPLICANT:
                                                                           APPLICANT:
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                                                                                                                                                                                                                                                2 LOVIGIOLPTP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: EP 97401404.5
FILING DATE: 18-JUN-1997
APPLICATION NUMBER: PCT/EP98/03307
FILING DATE: 03-JUN-1998
ATTORNEY/AGGNT INFORMATION:
NAME: YUN, Allen C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/467,008 FILING DATE: 20-Dec-1999 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                            ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: TRIEBEL, Frederic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                 Application US/08394442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 624 Ninth Street, N.W., Suite 300 CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 37.971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 422 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 202-628-5197
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                                                           Triebel,
                                                                                                                                                                                                                                                                                      Conservative
                                                                           Hercend, Thierry
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72.7%;
Lymphocytes, DNA Sequences Encoding These Proteins And Their Pharmaceutical And Biological Uses
                                   New Proteins Produced By Human
                                                           Frederic
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STREET: 4...
STREET: 4...
CITY: Washington

ADDRESSEE:

E: BROWDY AND NEIMARK, P.L.L.C. 419 Seventh Street, N.W., Suite 400

COUNTRY:

U.S.A.

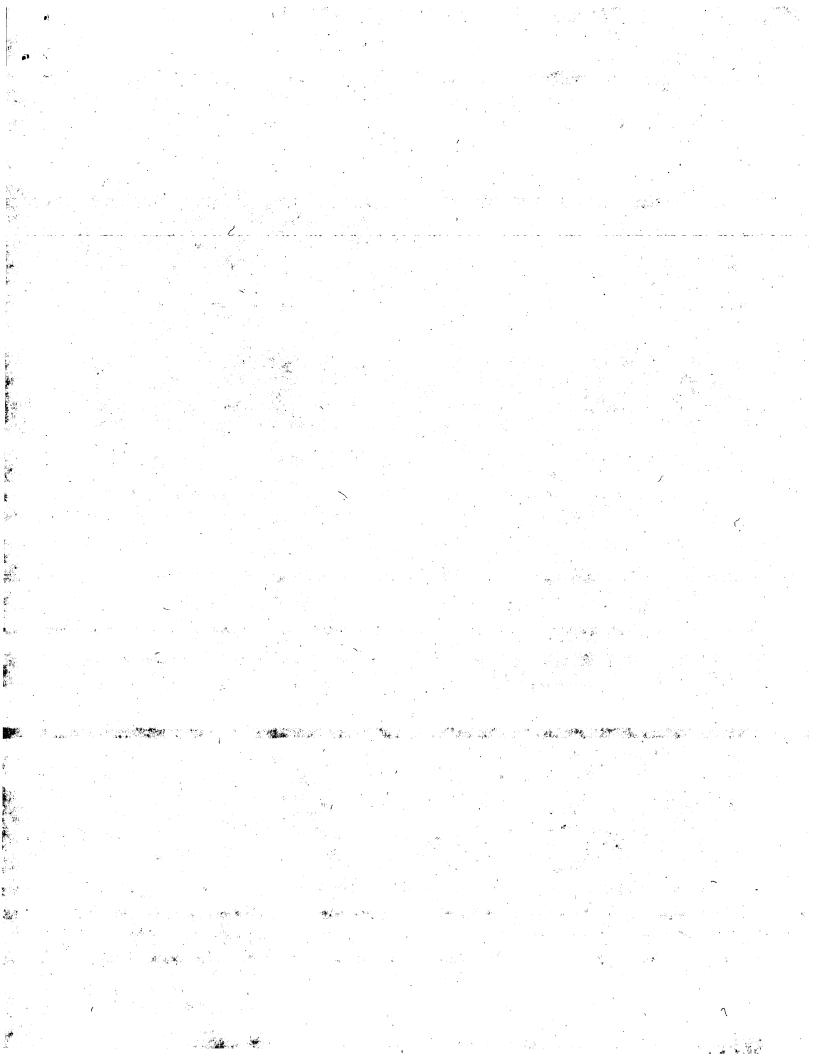
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Best Local Similarity 72.7%;
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: BROWDY, Roger L.
NAME: BROWDY, Roger L.
REGISTRATION UNMBER: 25,618
REFERENCE/DOCKET NUMBER: HERC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
TELEPHONE: (202) 737-3528
                                                       APPLICATION NUMBER: US 08
FILING DATE: 04-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                   APPLICATION NUMBER: US/08/474 988 FILING DATE: 07-UUN-1995 CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/416/478 APPLICATION NUMBER: US 08/416/478
                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TBM: PC-DOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
COPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS: LENGTH: 470 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/394,442
FILING DATE: 24-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICATION NUMBER: US 07/854,644 FILING DATE: 08-SEP-1992 TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        229 LTVLGLEPPTP 239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                                                                                                                             COUNTRY: U
ZIP: 20004
                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS
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                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                 3: BROWDY AND NEIMARK; P.L.L.C. 419 Seventh Street, N.W., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hercend, Thierry
Triebel, Frederic
                                                                                                                                                                                                                                                                                                                                                            U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS AND THEIR PHARMACEUTICAL AND BIOLOGICAL USES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NEW PROTEINS PRODUCED BY HUMAN
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Pred. No.
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
229 LTVLGLEPPTP 239
                                                                                                                                                         LENGTH: 470 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                         NAME: YUN, Allen C.
REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HERCEND-1B
                                                                                                                                             TOPOLOGY:
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              2 LQVLGLQLPTP 12
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                                                        Conservative
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202) 737-3528
                                                                    Score 39;
Pred. No.
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Search completed: June 4, 2001, 12:20:05 Job time: 518 sec



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Result
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Maximum Match 100%
Listing first 45 summaries
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length: 2000000000
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1: /cgn2_6/ptodata/1/paa/PCT_NEW_COMB.pep:*
2: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep::
3: /cgn2_6/ptodata/1/paa/US07_NEW_COMB.pep::
4: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep::
5: /cgn2_6/ptodata/1/paa/US08_NEW_COMB.pep::
6: /cgn2_6/ptodata/1/paa/US06_NEW_COMB.pep:
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PCT-US01-11988-2090
US-09-833-245-2090
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US-09-270-849B-180083
US-09-739-449-12942
PCT-US01-11988-1273
PCT-US01-11988-1274
PCT-US01-11988-1275
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1035, Ap
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PCT-US01-01339-4275	US-09-270-849B-184149	US-09-833-245-893	PCT-US01-11988-893	US-09-814-666-369	US-60-248-505-1177	US-09-628-359-17	US-60-248-505-797	US-09-270-849B-181560	US-60-248-505-1225	PCT-US01-04098A-3676	PCT-US01-04098A-1708	US-09-739-449-9005	US-09-813-872-4	US-09-739-449-12091	US-60-248-505-882	US-09-739-449-8808	US-09-833-245-1275
Sequence 4275, Ap	Sequence 184149,	Sequence 893, App	Sequence 893, App	Sequence 369, App	Sequence 1177, Ap	Sequence 17, Appl	Sequence 797, App	Sequence 181560,	Sequence 1225, Ap	Sequence 3676, Ap	Sequence 1708, Ap	Sequence 9005, Ap	Sequence 4, Appli	Sequence 12091, A	Sequence 882, App	Sequence 8808, Ap	Sequence 1275, Ap

### ALIGNMENTS

US-09-284-107-28

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Best Local S
Matches 12
                                                                                                              Sequence 1035, Application PC/TUS0101312
GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc., et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, a
FILE REFERENCE: PT232PCT
CURRENT APPLICATION NUMBER: PCT/US01/01312
CURRENT FILING DATE: 2001-05-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 28, Application US/09284107
GENERAL INFORMATION:
APPLICANT: Logtenberg, Ton
APPLICANT: de Kruif, Cornelis Adriaan John
TITLE OF INVENTION: METHODS AND MEANS FOR TITLE OF INVENTION: AND PROTEINS HAVING SPI
FILE REFERENCE: 313632000600
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1035
LENGTH: 141
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SEQ ID NO 28
LENGTH: 12
TYPE: PRT
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CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
PRIOR FILING DATE: 1997-10-07
PRIOR APPLICATION NUMBER: EP 96202791.8
PRIOR FILING DATE: 1996-10-08
                                                                      Prior application data removed - refer to PALM or NUMBER OF SEQ ID NOS: 1510
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Local Similarity 100.0%;
nes 12; Conservative (
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Pred. No. 0.00031;
); Mismatches 0;
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SPECIFIC AFFINITY FOR A TARGET
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                                                         OTHER INFORMATION:
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; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids.
PCT-US01-01339-2813
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                                                                                                                                                                      SEQ ID NO 1005
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Best Local Similarity
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Best Local :
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SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                 CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Human Genome Screwco, TITLE OF INVENTION: Nucleic Acids, Proteins, FITLE REFERENCE: PC006PCT
                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: PCT/US01/01329
                                                                                                                                                                                                                                                                                  equence 1005, Appli
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                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                   NERAL INFORMATION:
PPLICANT: Human Genome Sciences, Inc., et al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 10231
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                                                                                    NAME/KEY: SITE LOCATION: (1)
                                                                                                                  FEATURE:
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                                                                                                                                                          ENGTH: 110
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OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
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 Score 35;
Pred. No.
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Pred. No. 66;
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Best Local Similarity

Query Match

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                                                                                                                                            SEQ ID NO 2091
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OTHER INFORMATION: -09-833-245-2091
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                             PEATURE:
NAME/KEY:
LOCATION:
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                                                                                                                                                           SOFTWARE: PatentIn Ver.
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                                                                                                                                                                                               PRIOR FILING DATE:
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PRIOR APPLICATION NUMBER: 60/229, 358
PRIOR FILING DATE: 2000-04-12
PRIOR APPLICATION NUMBER: 60/256, 931
PRIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                  PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/229,
                                                                                    ORGANISM:
                                                                                                        TYPE: PRT
                                                                                                                                                                                                               RIOR APPLICATION NUMBER: 60/199, 384
                                                                                                                                                                                                                                RIOR APPLICATION NUMBER: 60/256, 931 RIOR FILING DATE: 2000-12-21
                                                                                                                                                                                                                                                                                                                                         'ILE REFERENCE: PF546PC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Human Genome Sciences, Inc
                                                                                                                                                                                                                                                                                                                                                    PPLICANT: Human Genome Sciences, Inc. ITLE OF INVENTION: Albumin Fusion Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: PCT/US01/11988
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                                                                                                                       ENGTH:
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                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/833,245 FILING DATE: 2001-04-12
                               SITE (87)
                                                                                    Homo sapiens
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           equals any of the naturally occurring L-amino acids
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PCT-US01-01339-3249
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; ORGANISM: Homo sapiens
PCT-US01-01339-3249
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                          Query Match
                                                                       NAME/KEY: SITE
LOCATION: (116)
OTHER INFORMATION: X
T-US01-11988-2090
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APPLICANT: Human Genome Sciences,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 10231
SOFTWARE: PatentIn Ver. 2.0
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ENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: PCT/US01/11988
CURRENT FILING DATE: 2001-01-12
                                                                                                                                                                                                                                                                                                                               RIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                     RIOR APPLICATION NUMBER: 60/229,
                                                                                                                               NAME/KEY: SITE
LOCATION: (109)
OTHER INFORMATION: Xaa
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                                                                                                                                                                                                                                                                                                                                            RIOR APPLICATION NUMBER:
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Local Similarity
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58.3%;
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             Score 34; DB 1;
Pred. No. 1.1e+02;
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SEQ ID NO 9638
LENGTH: 335
TYPE: PRT
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SEQ ID NO 2090
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APPLICANT: Human Ge
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LOCATION: (109)
OTHER INFORMATION: 1
NAME/KEY: SITE
LOCATION: (116)
                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
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CURRENT FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                               APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences
FILE REFERENCE: 38-10(15490)C
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CURRENT FILING DATE: 2001-04-12
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                                                                                                                     ORGANISM: Agrobacterium tumefaciens
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                                  Score 34;
Pred. No.
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Query Match
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5,5heq 6; Conserv:
                                                                                                 ; ORGANISM: Agrobacterium tumefaciens US-09-739-449-13150
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PCT-US01-04098A-1405
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SEQ ID NO 1405
LENGTH: 350
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                                                                                                                                                NUMBER OF SEQ ID NOS: 13351
EQ ID NO 13150
LENGTH: 843
                                                                                                                                                                                                   PRIOR FILING DATE:
                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
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CURRENT FILING DATE: 2000-12-19
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                                                                                                                                                                                                                                                                                                    APPLICANT: Hinkle, Gregory J. APPLICANT: Slater, Steven C.
                                                                                                                                      TYPE: PRT
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ERAL INFORMATION:
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APPLICATION NUMBER: 09/693,325
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FILING DATE: 2000-04-27
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55.6%;
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Pred. No., 3.2e
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               Mismatches
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                                             Length 843;
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; ORGANISM: Homo sapiens
US-09-809-391-328
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SEQ ID NO 328
LENGTH: 23
TYPE: PRT
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EARLIER FILING DATE: 1992-03-11 NUMBER OF SEQ ID NOS: 116
                                   EARLIER APPLICATION NUMBER: 07/964,589
EARLIER FILING DATE: 1992-10-21
EARLIER APPLICATION NUMBER: PV-709-92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: 186 Human Secreted proteins
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                                                                                                    FILING DATE: 1993-12-30
                                                                                                                 FILING DATE: 1994-06-15
APPLICATION NUMBER: 08/177,093
                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/486,756 FILING DATE: 1995-06-07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/178,115
                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/485,862
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Pred. No.
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; LENGTH: 257 ; TYPE: PRT ; ORGANISM: HUMAN US-09-178-115-51

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Prior application data removed - refer to PALM or file wrapper

NUMBER OF SEQ ID NOS: 1249

SEQ ID NO 1196

LENGTH: 334

LENGTH: 334

CRCANISM: Homo sapiens
FEATURE:
FEATURE:
LOCATION: (184)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
PCT-US01-01332-1196
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Search completed: June 4, 2001, 12:21:30 Job time: 592 sec
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GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc., et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: DZ020PCT
CURRENT APPLICATION NUMBER: PCT/US01/01332
CURRENT FILING DATE: 2001-05-09
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Matches 6; Conservative
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46 LELLGFQLP 54
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45 VKGINLPTP 53
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66.7%;
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GenCore version 4.5. Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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US-09-489-039A-7598
US-09-107-433-2896
US-09-450-969-6248
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PCT-US00-05882-1128
US-07-861-895-25
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Sequence 7770, Ap
Sequence 7770, Ap
Sequence 7598, Ap
Sequence 2896, Ap
Sequence 6248, Ap
Sequence 11281, A
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Sequence 1128, Ap
Sequence 25, Appl
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## ALIGNMENTS

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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-702-021-14
                                                        RESULT 2
PCT-US00-05882-1128
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Best Local Similarity
Matches 12; Conserv
           Sequence 1128, Application PC/TUS0005882
GENERAL INFORMATION:
APPLICANT: Craig Rosen,
                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/702,021
CURRENT FILING DATE: 2000-10-30
                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: FAIL MOLECULES AND USES THEREOF FILE REFERENCE: 7853-217
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                      289 VWEHVLEYLAVG 300
                                                                                                                                                     1 VWFHVLFYLAVG 12
Craig Rosen,
Steve Ruben
                                                                                                                                                                                                                                                                                                                                             FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                        C. Fraser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09702021
                                                                                                                                                                                    Conservative
                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                      0
                                                                                                                                                                                    Score 68; DB 21;
Pred. No. 0.0083;
Mismatches (
                                                                                                                                                                                                                     DB 21;
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                                                                                                                                                                                                                Length 374;
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FITLE OF INVENTION: Human Cancer Associated Gene

Sequences and Polypeptides

CURRENT APPLICATION NUMBER: PCT/US00/05882

REFERENCE: PA106PC

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; NAME/KEY: SITE
; LOCATION: (349)
; OTHER INFORMATION:
PCT-US00-05882-1128
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           equence 25, Application US/07861895 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EARLIER FILING DATE: 1999-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ARLIER APPLICATION NUMBER: 60/124,270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OCATION:
                                            REFERENCE/DOCKET NUMBER: 17
REFERENCE/DOCKET NUMBER: 17
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AME/KEY: SITE
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                                                                                                                                                                            FILING DATE: 25-OCT-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Rel
                                                                                                                              TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORRESPONDENCE ADDRESS
                                                                                           NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768.
                                                                                                                                                                                          APPLICATION NUMBER: PCT/A
                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                            OPERÁTING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INVENTION:
FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                        3000 K Street, N.W.,
                              (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HULETT, Mark D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HOGARTH, Phillip M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IERINO, Francesco L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Foley & Lardner
                                                                                                                                         UMBER: AU PJ 7045
25-OCT-1989
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Pred. No.
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Query Match
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Whiches 6; Conserv
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                                                                                                                                                            SEQ ID NO 7770
LENGTH: 370
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Best Local Similarity
Matches 10; Conserv
                                               Query Match
Best Local Similarity
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CTITLE OF INVENTION:

FILE REFERENCE: 107
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LENGTH: 370
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                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/252,691C CURRENT FILING DATE: 1999-02-18 PRIOR APPLICATION NUMBER: US 60/094,145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -09-252-691-7770
                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 1998-02-18 NUMBER OF SEQ ID NOS: 11326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/252,691B
CURRENT FILING DATE: 1999-02-18
NUMBER OF SEQ ID NOS: 11324
                                                                                                                                                                                                                                               PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
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2 WFHVLFYLA 10
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                                                                                                                                                                                                                                                                                                                                            Keith G. Weinstock et al. TENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROBACTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Keith G.
                                Conservative
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                                                                                                                                                                                                                                                                                                          N: CLOACAE FOR DIAGNOSTICS
                                                                                                                                                                                                                                            1998-07-24
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                                                                                                                                                                                                                                                    1999-02-18
IBER: US 60/094,145
                                                61.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61.8%;
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                                               Score 42;
Pred. No.
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Pred. No.
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131 WYHGIFYLA 139

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Query Match
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US-09-107-433-2896
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JS-09-489-039A-7598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2896, Application US/09107433
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
SEQ ID NO 7598
LENGTH: 410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/489,039/
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ITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA

ITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                           INFORMATION FOR SEQ ID NO: 2896:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                    APPLICATION NUMBER: 60/ 085131
FILING DATE: May 12, 1998
APPLICATION NUMBER: 60/051553
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/107,433 FILING DATE: 30-Jun-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (781)893-5007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
TYPE: amino acid
                                                                                                                                         REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-011
                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: <Unknown>
SOFTWARE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: <Unknown>
                                                                                                                                                                                 NAME: Ariniello, Pamela Deneke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: Massachusetts
                   LENGTH: 104 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ITY: Waltham
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BER: US 60/117,747
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45.5%;
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Pred. No.
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                                                                                                                  ORGANISM: Enterobacter cloacae US-09-252-691-11281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: S.epidermidis
US-09-450-969-6248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-450-969-6248
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/252,691B
CURRENT FILING DATE: 1999-02-18
NUMBER OF SEQ ID NOS: 11324
SEQ ID NO 11281
LENGTH: 418
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SEQ ID NO 6248
LENGTH: 244
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                                                                                                                                                                                                                                                                                                                    Sequence 11281, Application US/09252691B GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: PATH99-09A
CURRENT APPLICATION NUMBER: US/09/450,969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Lynn Doucette-Stamm et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCC
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 1999-11-29
                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                     FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                            139 IWFYLIFFLFFG 150
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2 WFHVLFYLAVG 12
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LOCATION: (B) LOCATION 1...104
SEQUENCE DESCRIPTION: SEQ ID NO: 2896:
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                                                                                                                                                                                                                                                                   Keith G. Weinstock et al.
VENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROBACTE
VENTION: CLOACAE FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/09450969
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Best Local Similarity 45...
5; Conservative
                              SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 35611
LENGTH: 239
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SEQ ID NO 11281
                                                                                    CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
                                                                                                                                ENERAL INFORMATION:

APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins
FILE REFERENCE: File Reference: 7326-094
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/417,507
CURRENT FILING DATE: 1999-10-14
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PRIOR FILING DATE: 1998-02-18
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ORGANISM: Drosophila melanogaster
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ITLE OF INVENTION:
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NUCLEIC ACID AND
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RESULT 15
US-09-741-669-400
; Sequence 400, Application US/09741669
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Best Local Similarity
Thes 7; Conserv
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US-09-417-507-28756
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CURRENT FILING DATE: 1999-10-14
NUMBER OF SEO ID NOS: 44312
SEQ ID NO 28756
                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 28756, Application US/09417507 GENERAL INFORMATION:
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SEQ ID NO 50828
LENGTH: 239
TYPE: PRT
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APPLICANT: Homburger et al
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                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACITITLE OF INVENTION: FUMIGATUS FOR DIAGNOSTICS FILE REFERENCE: PATH99-10
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KEITH G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094
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GENERAL INFORMATION:
APPLICANT: Forsyth, R. Allyn
APPLICANT: Ohlsen, Kari L.
APPLICANT: Ohlsen, Kari L.
APPLICANT: Oyskind, Judith W.
TITLE OF INVENTION: Genes identified as required for
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9245, Ap
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### ALIGNMENTS

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RESULT 2
US-09-739-449-9245
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SOFTWARE: Fasi
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/739,449
CURRENT FILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: US 09/514,000
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 13351
SEQ ID NO 9245
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CURRENT FILING DATE: 1999-10-25
PRIOR APPLICATION NUMBER: PCT/NL97/00557
                                                                                                                APPLICANT: Hinkle, Gregory J.
APPLICANT: Slater, Steven C.
TITLE OF INVENTION: Agrobacterium tumefaciens
FILE REFERENCE: 38-10(15490)C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: METHODS AND MEANS FOR TITLE OF INVENTION: AND PROTEINS HAVING SETILE REFERENCE: 313632000600
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APPLICANT: de Kruif, Corne
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US-09-270-849B-183288
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Best Local Similarity
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Best Local :
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SEQ ID NO 11606
LENGTH: 331
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Best Local Similarity
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LENGTH: 239
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 09/514,000 PRIOR FILING DATE: 2000-02-23
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CURRENT FILING DATE: 2000-12-19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Insect genome survey devices
                                                                                                                                               TYPE: PRT ORGANISM: Agrobacterium tumefaciens
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ITTLE OF INVENTION: Agrobacterium tumefaciens Genome Sequences
FILE REFERENCE: 38-10(15490)C
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224 WFHLPFYL 231
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Pred. No.
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Pred. No.
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Pred. No.
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Query Match
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NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 182343
LENGTH: 141
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Matches 6; Conserv
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SEQ ID NO 193851
LENGTH: 61
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                                                                                       CURRENT APPLICATION NUMBER: US/09/270,849B CURRENT FILING DATE: 1999-03-17
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
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TITLE OF INVENTION: Insect genome survey devices
                                                                                                                             FILE REFERENCE:
                                                                                                                                                TITLE OF INVENTION: Insect genome survey devices
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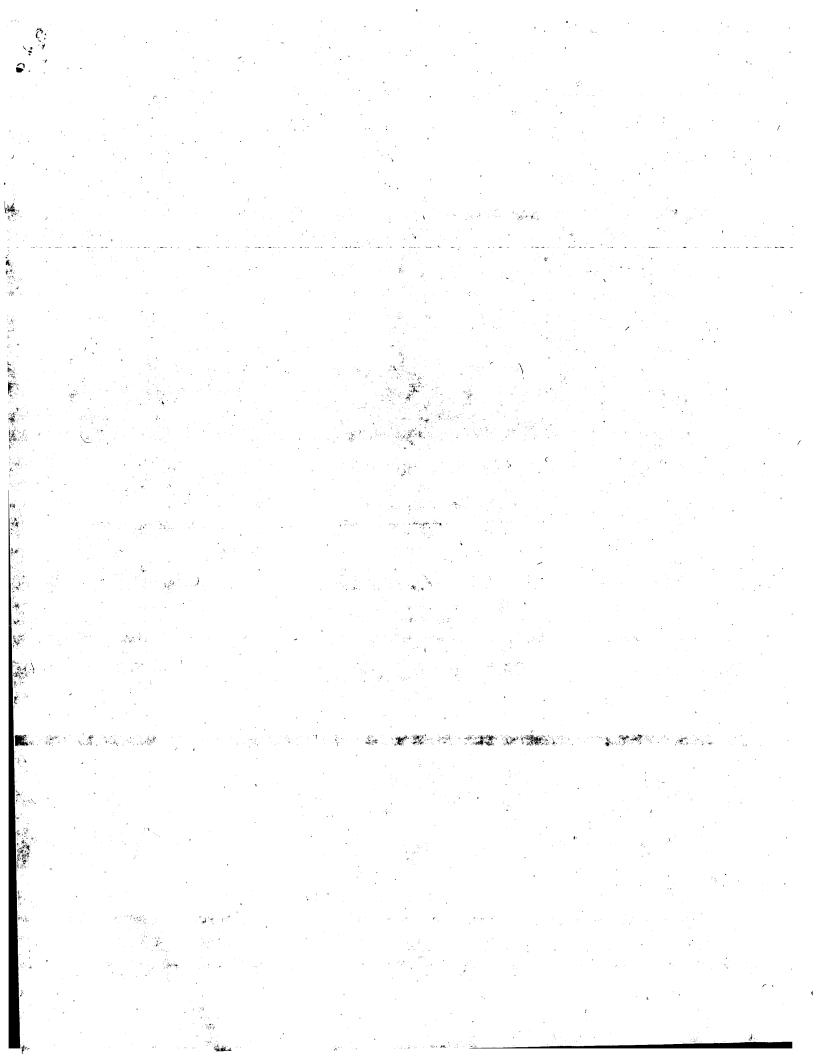
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; TYPE: PRT
; ORGANISM: Human
US-60-248-505-811
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Best Local Similarity
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; OTHER INFORMATION:
US-09-270-849B-182343
                                                                                                                                                                                                                                                                                                                                        RESULT 9
US-60-248-505-811
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Best Local Similarity
               Query Match
Best Local Similarity
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CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/60/248,505
CURRENT FILING DATE: 2000-11-15
                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                              NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                             FILE REFERENCE: cl000918
                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                             TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
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                                                                                                                                                                                                                                                                             FITLE OF INVENTION:
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
                                                                                                                        LENGTH: 1488
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RECEPTORS, NUCLEIC ACID MOLECULES ENCODING HUMAN GPCR
PROTEINS, AND USES THEREOF
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57.18;
             52.9%;
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71.4%;
               Score 36;
Pred. No.
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Pred. No. 3.8e+02
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   Mismatches
                DB 6;
5.7e+02;
                              Length 1488;
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Gaps
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Query Match
Best Local Similarity
Watches 4; Conserv
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; ORGANISM: Human
US-60-248-505-1121
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PCT-US01-01335-353
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PCT-US01-01339-5081
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SEQ ID NO 5081
LENGTH: 53
TYPE: PRT
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APPLICANT: Human Genome Sciences, Inc., et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Beasley,
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                                                                                                                    Sequence 353, Application PC/TUS0101335 GENERAL INFORMATION:
SEQ ID NO 353
LENGTH: 77
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CURRENT FILING DATE: 2001-03-17
NUMBER OF SEQ ID NOS: 10231
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CURRENT APPLICATION NUMBER: US/60/248,505
CURRENT FILING DATE: 2000-11-15
                                                            FILE REFERENCE: PC001PCT
CURRENT APPLICATION NUMBER: PCT/US01/01335
CURRENT FILING DATE: 2001-01-17
                          NUMBER OF SEQ ID NOS: 879
SOFTWARE: PatentIn Ver. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: PC006PCT
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                                                                                                                    TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
                                                                                                                                      APPLICANT: Human Genome Sciences, Inc., et al.
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24 IWFKIAFFLS 33
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Pred. No.
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; ORGANISM: Homo sapiens PCT-US01-01335-353

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Query Match
Best Local Similarity
Thehes 7; Conserv
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PCT-US01-04098A-1850
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Best Local Similarity 42...
6; Conservative
                  SEQ ID NO 187144
LENGTH: 123
TYPE: PRT
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SOFTWARE: Custom
SEQ ID NO 1850
LENGTH: 95
                                                                       CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                 APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens -US01-04098A-1850
ORGANISM: Artificial Sequence
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WIRDEN APPLICATION NUMBER: NOT YET ASSIGNED RICK FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 09/728,422
PRIOR APPLICATION NUMBER: 09/728,422
PRIOR FILING DATE: 2000-11-30
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TILE OF INVENTION: Novel Nucleic Acids and Polypeptides
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47 IWILPHLMLFLAVG 60
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FILING DATE: 2000-04-27
APPLICATION NUMBER: 09/496,914
FILING DATE: 2000-02-03
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NG DATE: 2000-09-01
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42.9%;
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0; Mismatches
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Pred. No.
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le+02;
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RESULT 15
US-09-270-849B-186130
Sequence 186130, Application US/09270849B
GENERAL INFORMATION:
APPLICANT: Swimmer et al.
TITLE OF INVENTION: Insect genome survey devices
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Best Local Similarity
7; Conserv
Search completed: June Job time: 592 sec
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; OTHER INFORMATION: Description
US-09-270-849B-187144
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                                                                                                             Query Match
Best Local Similarity bz..
Sches 5; Conservative
                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/270,849B
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 195450
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 186130
LENGTH: 62
                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                        TYPE: PRT
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Pred. No.
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                                                                                                                                                      1.1e+02;
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
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/SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:
/SIDS6/gcgdata/geneseq/geneseqp/AA2000.DAT:
                                                                6/gcgdata/geneseq/geneseqp/AA1994.DAT:
6/gcgdata/geneseq/geneseqp/AA1995.DAT:
6/gcgdata/geneseq/geneseqp/AA1996.DAT:
6/gcgdata/geneseq/geneseqp/AA1997.DAT:
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                                                                                                                                                                                                                                                                  /geneseqp/AA1988.DAT:
                                                                                                                                                                                                                                                                                                                                                            geneseqp/AA1984.DAT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (without alignments)
6.463 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         390729
                                           .DAT:
                                                                                                                                                        .DAT:
```

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

11	10	9	8	7	6	5	4	ω	2	1	No.	Result
55	55	55	55	. 55	55	55	55	55	55	55	Score	
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Match	Query
374	374	374	374	374	374	344	344	344	261	12	Match Length DB	
19	19	17	17	13	13	21	17	13	20	19	DB.	
W97833	W80448	W00859	R91438	R22550	R20811 ·	Y96183	R91439	R22549	Y33183	W60553	IJ	
Human Fc receptor	Human Fc receptor	Human FcRI (cDNA c	Human FcRI (cDNA c	Human macrophage-s	Human macrophage-s	Human macrophage-s	Human FcRI (cDNA c	Human macrophage-s	Human sFc-gammaRI	Oligopeptide from	Description	

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies

45	44	43	42		40			37								29				25	24	23	22	21	20	19	18	17	16	15	14	13	12
	32							33					33.	33	33	33	34	٠ 4	34	34	34	34	34	34	34	. 34	36	36	36	55	55	55	55
	58.2	8	8	8	0	0	0	0	0		0		0		60.0	.0	Ξ.	:-	61.8			:-		_			5	.51	65.5	00.		8	100.0
181	176	176	139	139	986	788	730	730	377	350	350	144	139	131	131	94	443	442	410	375	374	365	364	322	187	126	329	329	292	399	374	374	374
21	21	21	21	21									21	19	19	21	21	21	12	21	21	21	21	20	19	22	21	21	21	21	21	21	19
N	G38740	G20314	õ	G04555	W13670	W75919	W13669	P80618	W86195	R63143	P70645	G40794	G35305	•	W58851	53	~ 1	~1	R12428	G47110	G47083	G47111	G47084	W85718	W54052	в63956	Y82015	B28626	B28636	B43683	Y96226	Y96134	W97834 ·
																												•				-	
														٠.																			

## ALIGNMENTS

RESULT	ILT 1
MOUDDS	
ID	Wb0553 standard; peptide; 12 AA.
×	
AC.	· W60553;
×	
DΤ	18-AUG-1998 (first entry)
×	
Œ	Oligopeptide from extracellular domain of CD64.
×	•
ΚW	Extracellular domain; CD64; identification; antibody;
XW	immunohistochemical; immunofluorescent analysis; detection;
W.	cell transformation; mutation; anti; oncogene.
×	
<b>4</b> C	of inclination
PN	WO9815833-A1.
×	
PD	16-APR-1998.
PF	07-OCT-1997; 97WO-NL00557.
X	
PR	08-OCT-1996; 96EP-0202791.
PA X	(UYUT-) RIJKSUNIV UTRECHT.
×	
PI	De Kruif CA, Logtenberg T;
DR S	WPI: 1998-240964/21.
XX	
PT	Identifying peptide(s) binding specifically to protein target -
PT	surface and testing for binding to i
PŢ	oligo:peptide derived from the target, useful for, e.g. identify
ŢŢ	specific antibodies

method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or thousand differentiate between different forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode

affinity binding to the CD64-covered rods

domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were iso

English

W60537-61 are derived from the extracellular

The specification describes

isolated by describes a

Synthetic oligopeptides domain of CD64. They wer Example 1; Page 29; 40pp;

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XPXPX
Three-dimensional structures and models of Fc receptors, computer-assisted drug design
                                                                                                       Baell JB, I
                                                                                                                                                                                             11-SEP-1998;
06-FEB-1998;
                                                     WPI; 1999-539978/45
                                                                                                                                                                                                                                                    04-FEB-1999;
                                                                                                                                                                                                                                                                                                                        W09940117-A1
                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                          anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                            atomic coordinate; bloactive compound design; drug design; therapy; inhibitor; 'Fc-gammaR; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human sFc-gammaRI protein fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15~NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Y33183 standard; Protein;
                                                                                                                                                      (ILEX-) ILEXUS PTY LTD .-
                                                                                                                                                                                                                                                                                     12-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                           inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                            tissue damage; hypersensitivity; inflammatory cell recruitment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FC receptor; Fc-gammaRIIa; human; FCR; model; three-dimension; 3-D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N
                                                                                                       Epa V, Garrett TPJ, McKe
                                                                                                                                                                                                                                                                                                                                                                                                           modulator; Fc-gammaRIa;
                                                                                                                                                                                             98US-0099994
98US-0073972
                                                                                                                                                                                                                                                  99WO-IB00367.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         261
                                                                                                       TPJ, Hogarth PM, Matthews McKenzie IFC, Pietersz GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A
                                                                                                                                                                                                                                                                                                                                                                                                             Immune
                                                                                                                                                                                                                                                                                                                                                                                                                                          n; computer-assisted;
Fc-epsilonR; IgG; IgE;
                                                                                                                                                                                                                                                                                                                                                                                                       function regulation;
                                                                                                                       Matthews
                useful
```

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90

Example

10;

Page

94a;

160pp; English.

neoplasms,

etc.

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δÃ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIa and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, inchemical compounds can be designed to block binding of immunoglobulin to chemical compounds can be designed to block binding of immunoglobulin to
                                                                                                           New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                             WPI; 1992-056864/07
N-PSDB; Q21179.
                                                                                                                                                                                                                Seed
                                                                                                                                                                                                                                                                                13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotides, peptides, peptidomimetics and small organic moleculus by computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of the FCR can be used to design bloactive chemical compounds, e.g.
                                                                                                                                                                                                                                                 (GEHO-)
                                                                                                                                                                                                                                                                                                               15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                  23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                 WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                               Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human macrophage-specific FcRI receptor encoded by clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R22549;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R22549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 11; Page 316-317; 326pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 an Fc receptor protein using various computer programs and models.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   174 vlnasvtsplle 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 VLNASVTSPLLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    w
                                                                                                                                                                                                              В,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 Similarity
12; Conser
                                                                                                                                                                                                                                                 GEN HOSPITAL CORP
                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; llarity 100.0%; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                  90US-0553759
                                                                                                                                                                                                                                                                                                                 90WO-US04986.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12
                                                                                                                                                                                                                Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 55; DB 20;
Pred. No. 0.0036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                         it -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        molecules,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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밁 Qy

Matches Query Match

Local Similarity nes 12; Conser

100.0%; ilarity 100.0%; Conservative (

0

Mismatches

Score 55; DB 19; Pred. No. 6.2e-05;

Length 12; Indels

0;

Gaps

0

Sequence

12

ĀĀ,

by mutation in (anti)oncogenes.

antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused

are used, e.g.

peptides can be isolated and used for recombinant production of

antibodies.

88888888888888

0

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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
The amino acid sequence (R91439) of human FCRI was detd. from a cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence of the p98 product is truncated compared with those of the p135
                                                                                                                                                                                                                                     Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                             N-PSDB; T14718.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FcRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six
                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                                                                Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R91439 standard; Protein; 344 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain,
                                                                                                                                                                                          Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cell surface antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                           (GEHO ) GEN HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                  1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                Seed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      89US-0379076.
90US-0553759.
93US-0139273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92US-0983647
88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                8,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                             CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cloning; imm
ector; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid 58 is Leu in clone pl35 translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 55;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunoselection; immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.0052;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                           useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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B 8

Misc-difference

/note= 213

"encoded

by AAT"

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RESULT
Y96183
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         33
                                                                                          Ξ
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                                                                                                                         33
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            and p90 products (see also R91438 and W00859). if of FCRI and a receptor ligand will be helpful in potency of antibodies in therapy.
                                      Misc-difference
                                                            Misc-difference
                                                                                 Misc-difference
                                                                                                                         Misc-difference
                                                                                                                                             Misc-difference
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                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                             amyloidosis; multiple sclerosis; parasitic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                      panning; immunodiagnosis; diagnosis; immunotherapy; gene ty
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human macrophage-specific FcRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Y96183 standard;
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                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macrophage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              193 vinasvtspile
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nilarity 100.0%;
Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            FCRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
/note-
140
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117
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A fusion protein n increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 344;
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                                                                                                                                                                                                                                                                                                                                                                                                                             receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy;
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Misc-difference

"encoded"

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sclerosis. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
  surfaces other than macrophages, is useful protein of FcRI and a receptor ligand will the potencies of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                The present sequence is that of a human macrophage specific rcRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Stamenkovic I,
                                                                                                                                                                           immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in
                                                                                                                                                                                                                       CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
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                                                                                                                                                                                                                                                                                                              the invention designed to isolate CSA nucleic acids. The met
based upon transient expression of a CSA in eukaryotic cells
                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 10; Column 53-55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-586382/55.
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                                                                                                                                nimals, including humans. These disorders include asthma, mmune-complex disease, amyloidosis, parasitic diseases or multiple
                                                                                                                                                                                                                                                                                       physical selection of cells expressing the antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUL-1989;
23-MAR-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-OCT-1998;
                                                                                                                                                                                                                                                                  panning on) an antibody-coated substrate such as a culture dish.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-FEB-1988
                                                                                                                                                       including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0983647
88US-0160416
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220
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338
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305
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268
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216
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333
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ligand will therapy.
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                                                                                                                                                                                                                                                                                     otic cells and by adhesion to
                                                                                                                                                                                                                                                                                                                                   The method is
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Query Match 100.0%; Best Local Similarity 100.0%; Matches 12; Conservative (

0,

Mismatches

Score 55; DB 21; Pred. No. 0.0052;

DB 21;

Length 344; Indels

0;

Gaps

0;

21-MAY-1992 '(first entry)

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1 VLNASVTSPLLE 12

Sequence

344 AA;

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RESULT
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R20811
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                                                                                                                                                                                                                                                                                                                                                 This amino acid sequence was predicted from the cDNA sequence of C.DNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones predicts Val. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                  Query Match 100
Best Local Similarity 100
Matches 12; Conservative
                                                 R22550;
                                                                                                                                                                                                                                                                                                                    Sequence
                                                                              R22550 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R20811 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New CD53 cell surface antigen and DNA encoding it – for immuno-therapy and diagnosis of haematopoietic neoplasm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; Q21178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1992-056864/07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human macrophage-specific FcRI receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-MAY-1992
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                                                                                                                                                                                                                                                                                                                      374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                Protein; 374
                                                                                                                                                                   204
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                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                Score 55; DB 13; -Pred. No. 0.0058;
                                                                                AA.
                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               encoded by clone p135
                                                                                                                                                                                                                                                                     DB 13;
                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neoplasms,
                                                                                                                                                                                                                                                                Length 374;
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Human

macrophage-specific FcRI receptor encoded by clone p90

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  ETT FTT XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                             and clone p30 predicts a Thr residue. At position 58, p30 predicts val while clones p135 and p38/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FGRI and a receptor ligand will be belieffly the protein of FGRI and a receptor ligand will be
                                                                                                      Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                  Human
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90. It differs from the sequence predicted from cDNA clone p135 (see Q21178) at position 25; p135 encodes a Ser residue clone p135 (see Q21178) at position 25; p135 encodes a Ser residue
 Misc-difference
                                                                               Homo sapiens
                                                                                                                                                                          30-OCT-1996 (first entry)
                                                                                                                                                                                                     R91438
                                                                                                                                                                                                                                R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; Q21180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JÜL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                     helpful to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 94a; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-056864/07
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                                                                                                                                                                                                                                                                                                                            1 VLNASVTSPLLE 12
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                                                                                                                                               (cDNA clone p135 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                     increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                          374 AA;
                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90WO-US04986
                                     Location/Qualifiers
                         /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amiot M;
                                                                                                                                                                                                                                                                                                                                                                   100.0%;
           "amino acid 25 is Thr in clone p90 and p98
translated products"
                                                                                                                                                                                                                                                                                                                                                       0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English
                                                                                                                                                                                                                                                                                                                                                                   DB 13;
0.0058;
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PRR XXX
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W00859
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Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
 01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The amino acid sequence (R91438) of human FCRI was detd. from a cDNA clone, pl35 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p96 (T14718), coded for a FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                               25-FEB-1988
                                                                                             09-APR-1996
                                                                                                                             US5506126-A
                                                                                                                                                           Homo sapiens
                                                                                                                                                                                          therapy; diagnosis;
                                                                                                                                                                                                          Cell surface antigen; cloning; immunoselection; immunotherapy;
                                                                                                                                                                                                                                         Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                            30-OCT-1996
                                                                                                                                                                                                                                                                                                          W00859;
                                                                                                                                                                                                                                                                                                                                          W00859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Column 55-56; 79pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1996-200279/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                       193 vlnasvtsplle 204
                                                                                                                                                                                                                                                                                                                                                                                                                                      1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                        standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                          (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0983647.
88US-0160416.
89US-0379076.
89US-0553759.
93US-0139273.
                                                               88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vector; FcRI; Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                          374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 55;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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0

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RESULT
W80448
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Best Local Similarity
Matches 12; Conser
Misc-difference
                           Misc-difference
                                                      Misc-difference
                                                                                   Misc-difference
                                                                                                               Misc-difference
                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The amino acid sequence (W00859) of human FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cloning of cDNA encoding cell of diagnostic and therapeutic
                                                                                                                                        Misc-difference
                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                           Misc-difference 44
                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                        cloning
                                                                                                                                                                                                                                                                                                                                                                     Fc receptor I; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                               Human Fc receptor I.
                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                         W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W80448 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       193 vlnasvtsplle 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90US-0553759.
93US-0139273.
                           /note=
141
/note=
159
                                                      /note=
103
                                                                                   /note=
99
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85
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                                                                                                                                                                                                                                                      /note=
23
                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                     note=
                                                                                                                                                                                   'note=
                                                                                                                                                                                                             'note='
                                                                                                                                                                                                                                        note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
                                                                                                                                                                                                                                                                                                                                                             cell surface antigen; lymphocyte; human;
             "encoded
                                                                                                                                                                                                                                                                    "encoded by TGG"
                                                                   "encoded
                                                                                                                       "encoded by
                                       "encoded by
                                                                                                                                                                                                                                        "encoded by CTC"
                                                                                                                                                    "encoded by CCC"
                                                                                                                                                                                "encoded by CTG"
                                                                                                                                                                                                            "encoded by GAC"
                                                                                              "encoded by TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 55; DB 17;
Pred. No. 0.0058;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              surface antigen - useful for isolation
             Ьy
                                                                  bу
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                  CAA"
                                                                                                                           AAT"
                                         ccc"
              GGC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT -W97833 A X I

W97833;

W97833 standard;

Protein; 374

DЬ Qy

193 vlnasvtsplle 204

1 VLNASVTSPLLE 12

Matches

12;

Query Match
Best Local Similarity

100.0%; ilarity 100.0%; Conservative

0

Score 55; DB 19; Pred. No. 0.0058; Mismatches

Length Indels

374;

Gaps

0;

Sequence

374

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33333
                                                                                        the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                       immunotherapeutic applications, treatment of immune-mediated in
                                                                                                                                                                                                                                                                                           expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by
                                                                                                                                                                                                                                                                                                                               were also obtained.
                                                                                                                                                                                                                                                                                                                                                                                             cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                  This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone pl35 (see V63456) isolated using a rap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5830731-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                          in eukaryotic cells and selection of cells expressing adhesion to an antibody-coated substrate. The method
                                                                                                                                                                                                                                                                                                                                 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see x07372) and p98 (see x07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; V63456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1998-609251/51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                           immunoselection cloning method from a cDNA library expressed in opening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP
                                  including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97US-0861205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0983647
88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-0861205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          890S-0379076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=
256
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176
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171
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                                  humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "encoded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "encoded
                                                         infections
                                                                                                                                                                                                                                                                                                                                                                                                                                (see V63456) isolated using
                                                                         including the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     by GGG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bу
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bу
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAC"
                                                                                                                                                                                                                                                                          is useful
                                                                                                                                                                                                                                                                                                                                 from mammaliar
                                                         disorders
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in cos

for

07-JUN-1999

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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                 This is the amino acid sequence of human Fc receptor I (FCRI), a deduced from cDNA clone p90 (see x07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note=
Misc-difference 103
                                                                                                                                                New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                       WPI; 1998-609251/51.
N-PSDB; X07372.
                                                                                                                                                                                                                                                                                                                                                                                                                                          US5830731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note=
Misc-difference 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
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                                                                                                                      Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                     21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                03-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cloning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fc receptor I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human Fc receptor I.
              were also obtained
                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL
                                                                                                                                                                                                                                                                                                    21-MAY-1997
                            induction therapy
                                          trom a single patient undergoing extracorporeal
                                                     The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                               92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note=
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. 85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FCRI;
                                                                                                                                                                                                                                                                                                    97US-0861205
                                                                                                                                                                                                                                                                                                                                                                                     97US-0861205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note=
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                             Clones p135
                                                                                                                                                                                                                                                                         CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded by CTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "encoded by CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "encoded by TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "encoded by CTG"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "encoded by GGG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "encoded by CCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded by AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *encoded by
based on transient expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             γģ
                          (see V63456) and p98 (see X07373)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W97834 standard; Protein; 374
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Best Local Similarity 100.0%;
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       adhesion to an antibody coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W69440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
Macrophage; FcRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
                                                                                                            Human macrophage-specific FcRI.
                                                                                                                                                     19-DEC-2000
                                                                                                                                                                                                                                   Y96134 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
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25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Column 53-54; 75pp; English.
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N-PSDB; X07373.
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                                                                                                                                                                                                                                                                                                                                             193 vlnasvtsplle 204
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88US-0160416.
89US-0379076.
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97US-0861205
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25-FEB-1988;
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23-MAR-1990;
13-JUL-1990;
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Misc-difference 126
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Misc-difference 122
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Misc-difference 60
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Misc-difference 123
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                                                                                                                                      GGT"
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2000-586382/55

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RESULT 14
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Matches 12; Conser
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method o
                                                                                                                                                                                                                                                                                                     Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IGG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                     11-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        animals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
                                                                                                        03-DEC-1998;
                                                                                                                                         03-DEC-1998;
                                                                                                                                                                               07-JUN-2000
                                                                                                                                                                                                                   EP1006183-A1
                                                                                                                                                                                                                                                                                        systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Y96226 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immune complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         treatment of immune-mediated infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention designed to isolate CSA nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Column 53-55; 75pp; English.
                                WPI; 2000-367968/32
                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                 Human high affinity Fc receptor, FcgammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the potencies of antibodies in therapy.
                                                                  (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 VINASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vlnasvtsplle 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                        erythematosus; tumour.
                                                                                                        98EP-0122969
                                                                                                                                           98EP-0122969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                    FOERDERUNG WISSENSCHAFTEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 55;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diseases, and disorders in
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a novel method of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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glycosylation, cancer -
                                                                                                                                    Novel Fc receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
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Disclosure; Page 26-28; 60pp; English.

cells expressing Fc receptors. FrgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and esoinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant Fc receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immuneassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS, rheumatoid arthritis or MM. role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds The present sequence is the human high affinity Fc receptor, FcgammaRI. FcgammaRI is also known as CD64. Fc receptors play an important ç

Sequence 374

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Query Match
Best Local S
Matches 12
193 vlnasvtspile 204
                  1 VLNASVTSPLLE 12
                                  Similarity
12; Conserv
                                   Conservative
                                           100.0%;
                                   0;
                                           Score 55;
Pred. No.
                                   Mismatches
                                           DB 21;
                                                   Length 374;
                                   Indels
                                   0
                                   Gaps
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RESULT 08-FEB-2001 B43683 standard; Protein; 399 15 (first entry) Ā

Human cancer associated protein sequence SEQ ID NO:1128

naemostatic; thrombolytic; neurological disease; drug vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation; immune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; diagnosis; Human; cancer associated gene; cancer antigen; detection; dermatological; cytostatic; neuroprotective; thrombolytic; drug screening proliferative; vulnerary; coagulant; nootropic; immunomodulator; cancer;

Homo sapiens

XSX WO200055350-A1

08-MAR-2000; 2000WO-US05882

12-MAR-1999; 99US-0124270

PR XXX (HUMA-) HUMAN GENOME SCI INC

Rosen CA, Ruben

WPI; 2000-587533/55

N-PSDB; A27466

Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer -  $\!\!\!\!$ 

Claim 11; Page 1739-1740; 2352pp; English.

antithyroid; antiallergic; antibacterial; antiviral; dermatological; consumprotective; cardiant; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and coagulant; properties and solve preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosists and antagonists from the present conhibiting the proliferation, differentiation or mobilisation of immune cinhibiting the proliferation, differentiation on mobilisation of immune colls, to treat disorders of haematopoietic cells, autoimmune disorders, callergic reactions, graft versus host disease and organ rejection, condulate haemostatic or thrombolytic activity, modulate inflammation, condulate haemostatic or thrombolytic activity, modulate inflammation, concers, cardiovascular disorders, neurological disease and bacterial or collars infections. The peptides, nucleotides, antibodies, agonists and contraction, and contractions and contractions are perfectly and contraction. C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;

Sequence 399 AA;

Query Match 100.0%; Score 55; DB 21; Best Local Similarity 100.0%; Pred. No. 0.0063; Matches 12; Conservative 0; Mismatches 0 DB 21; Length 399; Indels 0; Gaps 0,

뫄 Š 1 VLNASVTSPLLE 12 ||||||||||| | 218 vlnasvtsplle 229

Search completed: June 4, 2001, 12:13:26 Job time: 199 sec

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Result
No.
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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Match
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1: /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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3.925 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                   Length
   1362
193
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780
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US-08-332-562A-94
US-08-764-100-27
US-08-721-488-8
5352575-7
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Sequence 12, Appl
Sequence 12, Appli
Sequence 6, Appli
Sequence 6, Appli
Sequence 1, Appli
Sequence 1, Appli
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Sequence 27, Appl
Sequence 8, Appli
Patent No. 5352575
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Sequence
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14, Appl
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3, Appl
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ALIGNMENTS	US-08-466-604-29 US-08-324-977-44 US-08-384-616-44 US-08-904-686A-44 US-08-9141-821-4 US-08-415-818-7 US-08-55-266A-14 PCT-US96-01444-7 US-08-554-987A-8 US-08-554-972-1 PCT-US91-08177-11 PCT-US91-08177-11 PCT-US91-08177-11 PCT-US91-08177-19 US-08-279-700-4 US-08-279-700-8 US-08-279-700-8 US-08-279-700-10
	Sequence 29, Appl Sequence 44, Appl Sequence 44, Appl Sequence 44, Appl Sequence 4, Appl Sequence 7, Appl Sequence 7, Appl Sequence 14, Appl Sequence 18, Appl Sequence 11, Appl Sequence 19, Appl Sequence 19, Appl Sequence 2, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 19, Appli Sequence 10, Appli Sequence 10, Appli Sequence 10, Appli

RESULT

US-				us ; ;
80 NI		0.		US-08-332-562A-94 ; Sequence 94, Appli ; Patent No. 5985599 ; GENERAL INFORMATI ; APPLICANT: MCK ; APPLICANT: HCG
FORN SEQUENTING TIME TIME TIME TIME TIME TIME TIME TIME	PRICE PRICE	COME CC	Appl Appl Appl SI	-332 lenc ent ent NERA NERA APPI
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TELEPHONE: ( TELEPHONE: () TELEX: 90413 ORMATION FOR S EQUENCE CHARAC LENGTH: 91 a TYPE: amino STRANDEDMESS: TOPOLOGY: 11 332-562A-94	FILING DATE: 3 FILING DATE: 3 FILING DATE: 3 FILOATION UM APPLICATION UM APPLICATION UM APPLICATION DATE: 2 FILING DATE: 2 ATTORNEY/AGENT ST REGISTRATION UM REGISTRATION UM REGISTRATION UM REFERENCE/DOCKE	CITY: Wash: STATE: D.C. COUNTRY: US ZIP: 20007- MPUTER READA MEDIUM TYPE: COMPUTER: I COMPUTER: I OPERATING SY SOFTWARE: P SOFTWARE: P	PLICANT: PLICANT: PLICANT: PLICANT: PLICANT: PLICANT TLE OF INV MBER OF SE RRESPONDEN ADDRESSEE: 3	562A-94 94, Ap 10. 5985 1NFORM CANT: CANT:
E: (20 (202) 904136 FOR SE FOR SE HARACT 91 am mino a mino a NESS: : lin	DATE: FICATION PLICATION PLICATION PATION PATE: DATE: DATE: BENT, BENT, RATIONUCE/DOCUMENT	Vashin D.C. USA )007-5 )007-5 READAB FYPE: IB SYPE: READAB	SCOUNT BC	94 Appl 9559 RMAI MC HC
NE: (202)672-5300 : (202)672-5399 904136 FOR SEQ ID NO: 9 CHARACTERISTICS: 91 amino acids amino acid DNESS: single Y: linear	ICAN NION	A BI BI BI STI STI	APPLICANT: HLDS, MAIGARE APPLICANT: SCOTT, Bernadet APPLICANT: BONADONNA, Lisa APPLICANT: HULETT, MAIK D. TITLE OF INVENTION: FC REC NUMBER OF SEQUENCES: 136 CORRESPONDENCE ADDRESS: ADDRESSEE: Foley & Lardn STREET: 3000 K Street, N	08-332-562A-94 equence 94, Application atent No. 5985599 GENERAL INFORMATION: APPLICANT: MCKENZIE, APPLICANT: HOGARTH, APPLICANT: HOGARTH,
) ID NO ERISTIC ino aci single	31-OCT 1994 ON: 435 ON: 435 ON DATA: ION DATA: US 0 27-MAY-1992 INFORMATION: Stephen A. INFORMATION: CKET NUMBER: 29, INFORMATION: TON INFORMATION	E FORM:  E FORM:  Floppy disk  I PC compati  EM: PC-DOS  EMI PC-DOS	, Ma ONNA ONNA ONNA N: M: N: DRES: ES: DRES Str	tior : ZIE,
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•	•	. 30	IMMUNOGLOBULIN	
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Query Match

61.8%; Score 34;

DB 2;

Length 91;

66.7%;

Pred. No.

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RESULT 3
US-08-721-488-8
Sequence 8, Application US/08721488
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Best Local :
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Sequence 27, App--
No. 5773700
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Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 17-MAR-1993
APPLICATION NUMBER: GB 9206016.9
FILING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO 5773700ris, Allen E.
REGISTRATION NUMBER: 34.490
REGISTRATION NUMBER: 34.490
                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 857-1125
NFORMATION FOR SEQ ID NO:
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2230 LNALAASPILE 2240
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                                                                                                                                                                                                                           MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 354-3592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ITLE OF INVENTION: Improvemen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JMBER OF SEQUENCES:
                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 06-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
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CITY: Palo Alto
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/214,064
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                                                                                                    2 LNASVTSPLLE 12
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                                                                                                                                                                                                                                                                                   3218 amino acids
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                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Goldbach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peters, Dirk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              van Grinsven J., Martinus Q.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sandoz Agro,
                                                                                                                                                                                                                                                         unknown
                                                                                                                                                  61.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robert W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US/08/764,100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US 08/032,235
                                                                                                                                                                                                                                                                                                                    27:
                                                                                                                                                                                                                                                                                                                                                                               137-1061
                                                                                                                                 Score 34; DB 1; Le
Pred. No. 8.7e+02;
1; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in or Relating to Organic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Version #1.25
                                                                                                                                                               Length 3218;
                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                      nt No. 5352575.

APPLICANT: PETROVSKIS, ERIK A.: POST, LEONARD E.; TIMMINS, JAMES G.
TITLE OF INVENTION: PSEUDORABIES VIRUS PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDITIM TYPE: Floppy disk
                                                                                                                                           FILING DATE: 20-APR-1990 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                            13 IFDAAVTXPTLE 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: 1:
                                            APPLICATION NUMBER: 784,787 FILING DATE: 04-OCT-1985
                                                                         APPLICATION NUMBER: 886,260 FILING DATE: 16-JUL-1986
                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                             APPLICATION NUMBER: 100,817 FILING DATE: 29-JUN-1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                            APPLICATION NUMBER: 801,799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                           1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ELEPHONE:
                                                                                                                                                                                           APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        131 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Genetics Institute,
87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treacy, Maurice
Spaulding, Vikki
Bowman, Michael
                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  McCoy, John
LaVallie, Edward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Merberg, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Racie, Lisa
NUMBER: 844,113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /) 450
876-5851
870: 8:
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                                                                                                                                                                                                                                                                                                                                                                                                                          60.0%;
50.0%;
                                                                                                                                                                            us/07/513,282
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                                                                                                                                                                                                                                                                                                                                                                                                                          Score 33;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 131;
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                                                                                                                                                                                                                                                                                                                                                                                                         0;
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Query Match
Best Local Similarity
"-+-hes 8; Conserve

Conservative

1; Mismatches Score 33; DB 6; Pred. No. 97;

60.0%; 72.7%;

Length 350; Indels

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5352575-7

LENGTH: 350

FILING DATE: 26-MAR-1986: SEQ ID NO:7:

В

260 VLNASVVSRVL 270

1 VLNASVTSPLL 11

US-08-572-225-1

equence 1,

Application US/08572225

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; STRANDEDNESS:
; TOPOLOGY: unk
; MOLECULE TYPE:
US-08-572-225-1
RESULT 6
US-08-232-538-12
                                                            В
                                                                                                                    Matches
                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                 TELEX: 66141 PENNIE NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tent No.
                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 25>227
REFERENCE/DOCKET NUMBER: 83
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-3660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
                                                            400 LNGSITSP 407
                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                       2 LNASVTSP 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10036
                                                                                                                                                                                                                                                       amino acid
                                                                                                                                                                                                                                                                                                                                                                                                        Halluin, Albert P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                     788 amino acids
                                                                                                                                                                                                                                                                                                                                415-854-3694
                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brenner, Mitch
                                                                                                                                                                                                                           unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sieron, Aleksander
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hojima,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prockop, Darwin J.
                                                                                                                                                                                                                                      unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RECOMBINANT C-PROTEINASE AND ITS USE FOR DRUG DEVELOPMENT FOR THE TREATMENT OF DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yoshio
                                                                                                                                    75.0%;
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                                                                                                                                                                                                                                                                                                                                                                              8389-031
                                                                                                                                                Score 33;
                                                                                                                                     Pred. No.
                                                                                                                      Mismatches
                                                                                                                                     5e+02;
                                                                                                                      1; Indels
                                                                                                                                             Length 788;
                                                                                                                      0;
                                                                                                                      Gaps
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CURRENT APPLICATION DATA:

OPERATING SYSTEM: SOFTWARE: Microso

Microsoft Word 6.0

DOS

COMPUTER:

IBM Compatible

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US-08-232-538-12
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                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                       Sequence 12,
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                                                                                                                                                                                                                          SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (908) 594-3905
TELEFAX: (908) 594-4720
INFORMATION FOR SEQ ID NO: 12:
                                                                                                                                                                                                                                          tent No.
ZIP: 07065-0900
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 661 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Wallen, John W.III
REGISTRATION NUMBER: 35,403
REFERENCE/DOCKET NUMBER: 18
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                        APPLICANT: THOMAS, I
                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                     223 VLNCTATTPL 232
                                                                                            ADDRESSEE: Me.
                                                                                STREET: P.O. CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 07065-0907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Rahway
                                     COUNTRY: USA
ZIP: 07065-0900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                  1 VLNASVTSPL 10
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                                                                                                                                                                                                                                          5861484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
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712380
                                                                                                                                                                                                                                                         Application US/08786164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P.O. Box 2000 126 E Lincoln Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Thomas,
                                                                                                                                                                                                           THOMAS, KENNETH A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                               Merck & Co., Inc.
O. Box 2000, 126 E. Lincoln Ave
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kenneth A.
                                                                                                                                                                                                                                                                                                                                                                                                                 58.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RICHARD L. RICHARD ENDOTHELIAL CELL
                                                                                                                                                                                              RICHARD
                                                                                                                                                              CELL GROWTH FACTOR
                                                                                                                                                                             INHIBITOR OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/232,538
                                                                                                                                                                                                                                                                                                                                                                                                 2; .Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18888IA
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 32; DB 1;
Pred. No. 3.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 661;
                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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PRIOR APPLICATION DATA:

CLASSIFICATION:

FILING DATE:

21-JAN-1997

US/08/786,164

APPLICATION NUMBER:

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Best Local Similarity
6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-232-538-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-786-164-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 equence 6, Application US/08232538, atent No. 5712380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NEORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                       TELEFAX: (908) 594-4720
NFORMATION FOR SEQ ID NO:
                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 687 amino acids
                                                                                                                                                                                                                                                CURRENT APPLICATION DATA: US/08/232,538
                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Wallen, John W.III
REGISTRATION NUMBER: 35,
                                                                                                                    REFERENCE/DOCKET NUMBER: 18
FELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 594-3905
                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 36,545
REFERENCE/DOCKET NUMBER: 18888DA
FELECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-3905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Mark Hand, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
 OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            223 VLNCTATTPL 232
                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: sir
                         STRANDEDNESS:
                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 07065-0907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Merck & Co., Inc.
STREET: P.O. Box 2000 126 E Lincoln Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 VLNASVTSPL 10
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                                                                                                                                                                                                                                                                                                                                                                                                        Rahway
                                           amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 661 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thomas, Kenneth A.
                linear
protein
                           single
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                                                                                                                                                                  35,403
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Pred. No.
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US-08-874-678-1

Sequence 1, Application US/08874678 Patent No. 5952199

GENERAL INFORMATION:

RESULT

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Query Match
Best Local Similarity
Ches 6; Conserv
                                         Query Match
Best Local Similarity
Trachas 6; Conserv
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                                                                                                                           ; MOLECULE TYPE: protein US-08-786-164-6
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                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 687 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-3905
                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: Mark Hand, J
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249 VLNCTATTPL 258
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                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: FILING DATE: 21-JA
                                                                                                                                                          TOPOLOGY:
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ZIP: 07065-0900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Merck & Co.,
                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                         1 VLNASVTSPL 10
                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Merck & Co., Inc.
P.O. Box 2000, 126 E. Lincoln Ave
                                                                                                                                                                                                                                                         908-594-4720
                                                        Conservative
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                                                                                                                                                          linear
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60.0%;
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INHIBITOR OF VASCULAR ENDOTHELIAL
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                                                                                                                                                                                                                                                                                                              36,545
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                                                                    Score 32;
Pred. No.
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Pred. No.
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; MOLECULE TYPE:
US-08-874-678-1
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NFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                           ent No.
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                  CORRESPONDENCE ADDRESS
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PRIOR APPLICATION DATA:
                                                                                                                                                   TITLE OF INVENTION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                  ITLE OF INVENTION:
IMBER OF SEQUENCES:
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TELEX: 910 277
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REGISTRATION NUMBER:
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                                                                STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United ZIP: 94111-4187
                                                  CITY: San
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STREET: Four Embarcadero Center, Suite 3400
                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                              1 VLNASVTSPL 10
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5. 6100071
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Y: United States
94111-4187
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1: United States
                              California
                                                              E: Flehr, Hohbach, Test, Albritton & Herbert Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     758 amino acids
                                                                                                                                                                                                       Presta, Leonard
                                                                                                                                                                                                                       Chen, Helen H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                   Ferrara, Napoleone
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                                                                                                                                                                                                                                       Davis-Smyth, Terri L
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                                              Francisco
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                                                                                                                                                   NOVEL INHIBITORS OF VASCULAR ENDOTHELIAL GROWTH FACTOR ACTIVITY, THEIR USES AND PROCESSES FOR THEIR
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60.0%;
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                                                                                                                                   PRODUCTION
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Pred. No. 3.8e+02;
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Best Local Similarity · 60.0%;
Matches 6; Conservative
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TELEX: 910 27729
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-232-538-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                   TELEFAX: (908) 594-4720 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (908) 594-3905
                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Wallen, John W.III
REGISTRATION NUMBER: 35,403
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LENGTH: 758 amino acid
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TELECOMMUNICATION INFORMATION
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ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
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                 SEQUENCE CHARACTERISTICS:
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ZIP: 07065-0907
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STRANDEDNESS: un
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                                                                TELEPHONE:
                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                       CITY: Rahway
                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Merck & Co., Inc.
STREET: P.O. Box 2000 126 E Lincoln Avenue
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10 277299
780 amino acids
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IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            GROWTH FACTOR
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INHIBITOR OF VASCULAR ENDOTHELIAL CELL
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                                   14:
                                                                                                     18888IA
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STRANDEDNESS: single

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US-08-786-164-14
                                                                             Query Match
Best Local Similarity
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Best Local Similarity
Matches 6; Conserv
                                                                Matches
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Patent No. 5861484
                                                                                                                                                                                                                                                     NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENERAL INFORMATION:
                                                                                                                                                   STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 780 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: THOMAS, KENNETH A. APPLICANT: KENDALL, RICHARD L. TITLE OF INVENTION: INHIBITOR OF VASCULAR ENDOTHELIAL TITLE OF INVENTION: CELL GROWTH, FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
249 VLNCTATTPL 258
                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 36,545
REFERENCE/DOCKET NUMBER: 188
CELECOMMUNICATION INFORMATION:
TELEPHONE: 908-594-3905
                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               249 VINCTATTPL 258
                                1 VLNASVTSPL 10
                                                                                                                                                                                                                                                                                                                                                                              TTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Microsoft Word 6.0 URRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 07065-0900
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 21-JA
                                                                                                                                                                                                                                                                                   TELEFAX: 908-594-4720
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 VLNASVTSPL 10
                                                               6;
                                                                                                                                                                                                       amino acid
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                                                                                                                                                                                                                                                                                                                                                                   Mark Hand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08786164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Merck & Co., Inc.
P.O. Box 2000, 126 E. Lincoln Ave.
                                                                                                                                                                                                                    780 amino acids
                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-JAN-1997
                                                                            58.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           58.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .US/08/786,164
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                                                                            Score 32; 1
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 32;
Pred. No.
                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9e+02;
                                                                                          Length 780;
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                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                             0,
                                                            Gaps
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RESULT 14 US-08-340-011-5

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US-08-901-710-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; MOLECULE TYPE: US-08-340-011-5
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 6
                                                                                                                                                                                                                                                                                          Sequence 5;
                                                                                                                                                                                                                                                                         Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 312/474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                          GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                            NUMBER OF SEQUENCES:
                                                                                                                          TITLE OF
                                                                                                                                                            APPLICANT:
                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                             TITLE OF INVENTION:
                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 36,989
REFERENCE/DOCKET NUMBER: 322
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 09-OCT-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                      256 VINCTATTPL 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                              STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 312/474-0448
COUNTRY: United States of America
                                                               ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                  1 VLNASVTSPL 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/9
FILING DATE: 09-OCT-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                                                                                                                                                                                                                                          INFORMATION:
              Chicago
: Illinois
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60606-6402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Meyers, Thomas C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chicago
                                                                                                                         T: Kaipainen, Arja
T: Matikainen, Marja-Terttu
INVENTION: FLT4, A RECEPTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                        Application US/08901710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08340011
                                              E: Marshall; O'Toole; Gerstein, Murray & Borun 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: Marshall, O'Too
6300 Sears Tower,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1311 amino acids
                                                                                                                                                                                                     Aprelikova, Olga
Pajusola, Katri
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                         Korhonen, Jaana
                                                                                                                                                                                                                                           Alitalo,
                                                                                                                                                                                           Armstrong,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide
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                                                                                                             THEREOF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07/959,951
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Power, 233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 32; DB 1;
Pred. No. 7.2e+02;
                                                                                                                            RECEPTOR TYROSINE KINASE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gerstein, Murray & Borun
South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 1311;
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                                                                                                                            AND USES
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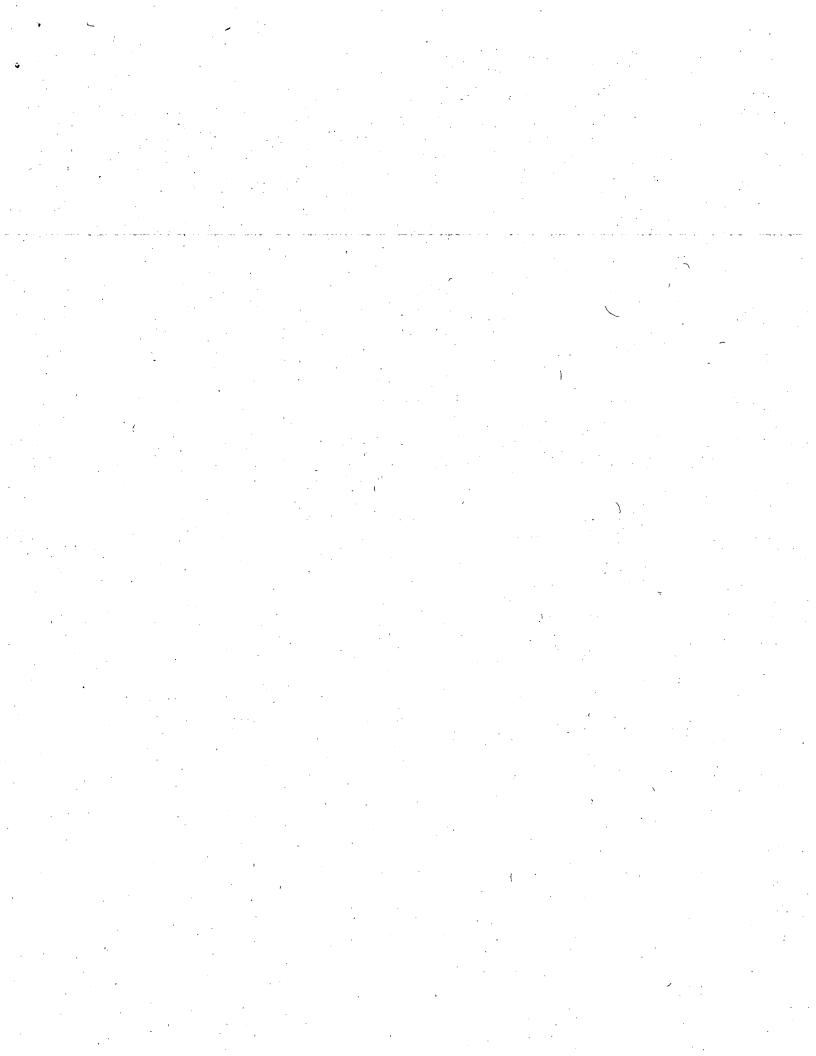
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; STRANDEDNESS:
; TOPOLOGY: line
; MOLECULE TYPE: p
US-08-901-710-5
Search completed: June 4, 2001, 12:14:28 Job time: 221 sec
                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION PC-DOS/MS-DOS
SOFTWARE: PATENTION: 435
CURRENT APPLICATION NUMBER: US/08/901,710
FILING DATE:
CLASSIFICATION NUMBER: US/08/901,710
FILING DATE:
CLASSIFICATION NUMBER: US/08/901,710
FILING DATE: 14-NOV-1994
PRIOR APPLICATION NUMBER: 08/340,011
FILING DATE: 14-NOV-1994
PRIOR APPLICATION NUMBER: 08/257,754
FILING DATE: 09-JUL-1994
PRIOR APPLICATION NUMBER: 07/959,951
FILING DATE: 09-OCT-1992
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 28113/33824
FELECOMMUNICATION INFORMATION:
TELEPHONE: 312/474-6300
TELEFAX: 312/474-0448
TELEX: 25-3856
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1311-amino acidd
STRANDEDNESS: single
TOPOLOCY: linear
                                                                                                                                                                                     Query Match 58.2%; Score 32; DB 3; Best Local Similarity 60.0%; Pred. No. 7.2e+02; Matches 6; Conservative 2; Mismatches 2
                                                                                                        256 VLNCTATTPL 265
                                                                                                                                                  1 VLNASVTSPL 10
                                                                                                                                                                                                                                                                                                                    peptide
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                  Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scoring table:
                                                                                                                                                                                                                                                                                                                                                     Database
                                                                                                                                                                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                            Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perfect score:
Pred. No. is the number of results predicted by chance score greater than or equal to the score of the result and is derived by analysis of the total score distribut
                                                                                                                                                                                                                                             Score
                                                                                                                                                                                                                                             Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                           BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-284-107-21
55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  June 4, 2001, 12:11:07; Search time 65.22 Seconds (without alignments)
12.645 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                     PIR_67:*
                                                                                                                                                                                                                                                                                                                                                                                                                                           198801 seqs, 68722935 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 VLNASVTSPLLE 12
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pir2:*
pir3:*
pir4:*
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	. 30
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60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	60.0	. 60.0
1082	991	986	823	730	615	524	524	430	428	418	405	379	350	288	251
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T43990	I49540	B58788	A58788	BMHU1	S77332 .	A82580	T06021	S77393	S46932	н82209	G70829	JC4289	VGBE63	T41112	A75482
hypothetical prote	procollagen C-endo	procollagen C-endo	procollagen C-endo	procollagen C-endo	NADH dehydrogenase	polyvinylalcohol d	hypothetical prote	cell division prot	hypothetical cell-	conserved hypothet	probable molybdenu	cell division prot	glycoprotein gp63	hypothetical prote	oxidoreductase, sh

## ALIGNMENTS

	RESULT 1
	A41357 Fc gamma (IgG) receptor I (high affinity) form b – human
	(man)
	C;Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000 C;Accession: A41357; S03019 P.Allen T.M. Scool B
	Science 243, 378-381, 1989 Science 243, 378-381, 1989 A;Title: Isolation and expression of functional high-affinity Fc receptor complementa
	A; Reference number: A41357; MUID:89100284 A; Accession: A41357
	A; Molecule type: mRNA A; Residues: 1-344 <all1></all1>
to have a being printed,	A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 R;Allen, J.M.; Seed, B.
icion.	Nucleic Acids Res. 16, 11824, 1988  A;Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (
	A; Reference number: SU3UI8; MUID:89U98339 A; Accession: SU3U18; MUID:89U98339
Description	A; Molecule type: mRNA A; Residues: 1-344 <all2></all2>
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334
c gamma (IgG) rec	A; Note: the authors translated the codon ACT for residue 25 as Ala
lcohol dehydrogen	C; Reywords: alternative splicing; immunoglobulin receptor; transmembrane protein
nypothetical glyci	F;117-170/Domain: immunoglobulin homology <imm></imm>
probable DNA-direc	100 09. 60070 55.
lcohol dehydrogen	cal Similarity 100.0%; Pred. No. 0.0038;
inkyrin-like prote	marches it; conservative o; mismarches o; indets o; edbs o;
ranslation elonga	Qy 1 VLNASVTSPLLE 12
HO85 protein - ye Drobable enoyl-CoA	Db 193 VLNASVTSPLLE 204
of gamma (IgG) rec	
cell division prot	A39878
ypothetical prote	FC gamma (IgG) receptor I-A (high affinity) precursor - human
rc gamma (IgG) rec	N;Alternate names: CD64 C:SDecles: Homo sapiens (man)
NA-export mediato	C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000
hosphomannomutase	C; Accession: A39878; I70304; B41357; S03018; I57525
nypothetical prote	T. Palol. Chem. 266, 13449-13455, 1991
hypothetical prote	A; Title: Gene organization of the human high affinity receptor for IgG, FcgammaRI (CD
	A; Reference number: A39878; MUID:91302383
lypothetical prote	A; Molecule type: DNA
conserved hypothet	A; Residues: 1-374 <van></van>

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submitted to the EMBL Data Library, January 1995
A;Description: Cloning and overexpression in Escherichia
A;Reference number: S51120
                                                           A; Accession:
                                                                                                                                     R; Cannio,
                       A; Residues: 1-347 <CAN>
                                     A; Molecule type: DNA
                                                                                                                                                     Accession: S51120
                                                                                                                                                                 Species: Sulfolobus sp.
Date: 19-Mar-1997 #sequence_revision 09-May-1997 #text_change 11-Jun-1999;
                                                                                                                                                                                                             lcohol dehydrogenase (EC 1.1.1.1) - Sulfolobus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rèsidues: 1-24,'S',26-337,'T',339-374 <AL2>
Cross-references: EMBL:X14356; NID 13131; PIDN:CAA32537.1; PI
Perez, C.; Wietzerbin, J.; Benech, P.D.
11. Cell. Biol. 13, 2182-2192, 1993
Title: Two cis-DNA elements involved in myeloid-cell-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cross-references: GB:M63830; GB:M63835; NID:g180277; Porges, A.J.; Redecha, P.B.; Doebele, R.; Pan, L.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Status: translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Accession: I70304
 Cross-references: EMBL:Z47543;
                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Reference number: I57525; MUID:93204964 Accession: I57525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title: Isolation and expression of functional high-affinity Reference number: A41357; MUID:89100284 Accession: B41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cross-references: GB:L03418; Allen, J.M.; Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Residues: 1-24,'S',26-337,'T',339-374 <AL1>
Cross-references: GB:X14356; GB:M21091; NID:g31331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Residues: 1-374 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    293-313/Domain: transmembrane #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GDB:135911; OMIM:146760
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: DNA
Residues: 1-8 <RE2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Molecule type: mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6-292/Domain: extracellular #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      L7-170/Domain: immunoglobulin homology <IMM2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      leic Acids Res. 16,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ference number: S03018; N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cession: S03018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                78,152,159,163,195,240/Binding
                                                                                                                                                                                                                                                                                                            193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /Domain: signal sequence #status
                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                              1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        iy: rc gamma receptor I; immunoglobulin homology
alternative splicing; glycoprotein: immunoglobu;
                                                                                                                                                                                                                                                                                                                                                                                    l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            378-381,
                                                           S51120
                                                                                                                                   Fiorentino, G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fc gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1q21-1q21
                                                                                                                                                                                                                                                                                                                                                                                100.0%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90, 2102-2109,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GB:S57204; NID:g298692; PIDN:AAD13887.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             155577;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11824, 1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                e of three cDNAs
MUID:89098339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MUID: 93055454
                                                                                                                                   Carpinelli, P.; Rossi, M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NID: g184840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1992
NID: 9623347;
                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                  Score 55; DB Pred. No. 0.0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 family
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                carbohydrate:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PIDN: AAA36049.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for the human high affinity Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                        0.0042;
                                                                                                                                                                                                                                                                                                                                                                                                                        DВ
   PIDN:CAA87591.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <EXT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                      Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PIDN:AAA35678.1; PID:g180279 Salmon, J.E.; Kimberly, R.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Asn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 in
                                                                                                                                     Bartolucci,
                                                                                                 coli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (covalent)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PID:g4261587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PID:g292169
                                                                                               of the genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PID:g31332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fc receptor complementary
   PID:g623348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptor;
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                                                                                                 NAD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Description:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Complex: homodimer
                                                                                                                                                                                                               R;Smith,
                                                                                                                                                                                                                                     C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nature 393,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rajandream, M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Pathway:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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F;174-203/Region: beta-alpha-beta NAD nucleotide-binding fold F;11,213/Modified site: N6-methyllysine (Lys) #status predicted F;38,68,154/Binding site: zinc, catalytic (Cys, His, Cys) #status
                                                                                                                                                                                                                   C;Keywords: alcohol metabolism;
F;23-337/Domain: long-chain alc
                                                                                                                                                                                                                                                    C;Superfamily: alcohol
                                                                                                                                                 38,68,154/Binding site;98,101,104,112/Binding
                                                                             Query Match
Best Local S
220 VINASVQDPLAE
                                                                                                                                                                                                                   Domain: long-chain alcohol dehydrogenase
                               VLNASVTSPLLE
                                                                                                                                                                                                                                                                     alconor
                                                                Similarity 66. 8; Conservative
                                                                                                                                                                                                                                                                                    catalyzes
                                                                                                                                                                                                                                                                 degradation
                                                                                                                                                 site: zinc, cata
nding site: zinc,
231
                                12
                                                                                                                                                                                                                                                    dehydrogenase; long-chain alcohol dehydrogenase
                                                                                 69.1%;
66.7%;
                                                                                                                                                                                                                                  heat-stable protein;
                                                                                 Score 38; DB Pred. No. 9.2;
                                                              Mismatches
                                                                                                                                                 noncatalytic
                                                                                                                                                                                                                                                                                    of primary
                                                                                                1;
                                                                                                                                                                                                                   homology
                                                                                                                                                                                                                                                                                    and secondary alcohols
                                                                                                                                                    (Glu,
                                                                                                Length 347
                                                                                                                                                    cys,
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                                                                                                                                                    Cys,
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Cys) #sta
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                                                                   0
                                                                                                                                                                                                                                    methy
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A; Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; A; Title: Deciphering the biology of Mycobacterium A; Reference number: A70500; MUID:98295987
                                                                                                                                                                                                                                                                                                                                                                                                                         hypothetical glycine-rich protein Rv0742 - C; Species: Mycobacterium tuberculosis
                                                                                                                                                           A; Experimental source:
                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 1-175 <COL>
                                                                                                                                                                                                                                                                                                                                                                                                          C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 22-Oct-199
                                                                                                                                                                            A; Cross-references: GB: AL021958;
                                                                                                                                                                                                                                                  A; Accession: A70824
                                                                                                                                                                                                                                                                                                                                                                                          C; Accession: A70824
                                                                                                                                                                                                                            A; Status: preliminary; nucleic acid sequence
4 ASVTSPLLE 12
                                                                                                                                                                                                                                                                                                                     537-544
                                 Similarity 7; Conser
                                                                                                                                                                                                                                                                                                                                                        Davies,
                                                                                                                                                                                                                                                                                                                                                                       Brosch,
                                 Conservative
                                                                                                                                                                                                                                                                                                                                  Rogers, J.; Rutter,
                                                                                                                                                                                                                                                                                                                                                      R.; Parkhill, J.; Garnier, T.; Churcher, C
                                                                                                                                                           strain
                                                  67.3%;
77.8%;
                                                                                                                                                           58; GB:AL123456; NID:g3261536; H37Rv
                                                  Score 37;
Pred. No.
                                                                                                                                                                                                                                                                                                                                        Seeger,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Mycobacterium tuberculosis (strain H37RV)
                                                  6.5;
                                                                                                                                                                                                                                not shown;
                                                                    2;
                                                                                                                                                                                                                                                                                  Whitehead, S.; Barrell, tuberculosis from the co
                                                                                                                                                                                                                                translation
                                                                                                                                                                              PIDN:CAA17509.1;
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                                                                                                                                                                                                                                    not shown
                                     Gaps
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                                                                                                                                                                            PID:e125
                                                                                                                                                                                                                                                                                                                                                          Holroyd
                                                                                                                                                                                                                                                                                                                                                                             Gordon
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A; Molecule type: DNA A; Residues: 1-361 < MTH>
                                                                                                      A; Reference number: A; Accession: D69005
                                                                                                                               A; Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta A; Reference number: A69000; MUID: 98037514
                                                                                                                                                                                                                                                                                                                   C;Species: Methanobacterium thermoautotrophicum
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000
A;Cross-references: GB:AE000876; GB:AE000666; NID:g2622140;
                                                                       A; Status: preliminary; nucleic acid sequence
                                                                                                                                                                                                                                                                                                                                                                    nypothetical protein MTH1037 - Methanobacterium thermoautotrophicum (strain Delta
                                                                                                                                                     . Bacteriol. 179, 7135-7155, 1997
                                                                                                                                                                                                                                Qiu, D.; Spadafora,
                                                                                                                                                                                                                                                              D.R.;
                                                                                                                                                                                                                                                                                            D69005
                                                                                                                                                                                                                                        Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; lafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski,
                                                                                                                                                                                                            R.; Vicaire, R.; Wang, Y.; Wi
Daniels, C.J.; Mao, J.; Rice,
                                                                             not shown; translation
                                                                                                                                                                                                               P.;
                                                                                                                                                                                                               Noelling,
  PIDN: AAB85533.1;
                                                                                                                                                                                                                                     Dubois, J.; Aldredge, T
J.; Gibson, R.; Jiwani,
                                                                                  not shown
                                                                                                                                                                                                                     Reeve,
  PID: g262
                                                                                                                                                                Η:
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source:

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R;Blattner, .... Mau, B.; Shao, X.
A.; Rose, D.J.; Mau, B.; Shao, X.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of F
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probable DNA-directed DNA polymerase (EC 2.7.7.7) -
C;Species: mitochondrion Pichia kluyveri
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #
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C; Superfamily: hypothetical protein MTH778
                                                                                                                      A;Molecule type: DNA
A;Residues: 1-12 <QUA>
R;Blattner, F.R.; Plun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-1118 <NOS>
A;Cross-references: EMBL:Y11606; NID:e1033579; PID:e321611; PIDN:CAA72340.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Reference number: A; Accession: T28426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             submitted to the EMBL Data Library, March 1997
A;Reference number: Z20485
                                A;Status: nucleic
                                            A; Accession: A64809
                                                                                                                                                                                                             FEBS Lett. 356, 183-187, 1994
                                                                                                                                                                                                                                                                                                                                        A; Title: Overexpression
                                                                                                                                                                                                                                                                                                                                                        Biochem. J.
                                                                                                                                                                                                                                                                                                                                                                                                       C; Date:
                                                                                                                                                                                                                                                                                                                                                                                                                                  N; Alternate names:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription regulator fark,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R; Nosek, J.
                                                                                                                                                                          Reference number: S50896; MUID:95104410
Accession: S50896
Status ----
                                                                                                                                                                                                                                                                                                                                                    Buck, D.; Guest, J.R. cochem. J. 260, 737-747,
                                                                                                                                                                     Status: preliminary
                                                                                                                                                                                                                                                Quail, M.A.;
                                                                                                                                                                                                                                                                             Residues: 1-240 <B
                                                                                                                                                                                                                                                                                                                            Reference number:
                                                                                                                                                                                                                                                                                                                                                                                       Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                    Species:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Keywords:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Accession:
                                                                                                                                                                                                                                               Cross-references: EMBL:X15790; NID:g41518; Quail, M.A.; Dempsey, C.E.; Guest, J.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                          Accession: S04645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              551
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                    10-Sep-1999 #sequence_revision 10-Sep-1999 sion: S04645; S50896; A64809; PC4138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 VLNASVTSPLL 11
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                                                                                                                                                                                                                                                                                                                                                                                                                    e names: gp30 pro
Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mitochondrion;
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                                                                                                                                                                                                                                                                           1-240 <BUC>
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                                                                                                                                                                                                                              Dempsey, C
, 183-187,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                              acid sequence not shown; translation not shown
                                                                                                           Plunkett III, G.;
Mau, B.; Shao, Y.
                                                                                                                                                                                                                                                                                                                            S04645;
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71.48;
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72.78;
                                                                                                                                                                                                                                                                                                                                                        1989
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                                                                                                                                                                                                                                                                                                                            site-directed MUID:89350876
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                                                                             Escherichia
                                                                                                                                                                                                                                                                                                                                        mutagenesis
                                                                                                                      C.A.;
                                                                                                                                                                                                                                                             PIDN:CAA33791.1;
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                                                                                                                         Perna,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #text_change 15-Oct-1999
                                                                                                                                                                                                                  regulator (FarR) in
                                                                                                                                                                                                                                                                                                                                                                                                    #text_change 10-Sep-1999
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                                                                             coli
                                                                                                                                                                                                                                                                                                                                          of.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Escherichia
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                                                                                                                        Z. T. ;
                                                                                                                                                                                                                                                                                                                                        the
                                                                                                                                                                                                                                                                                                                                        succinyl-CoA synthetase
                                                                                                                         Burland,
                                                                                                                                                                                                                                                             PID:g41519
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                                                                                                                      ٧.;
                                                                                                                                                                                                                Escherichia coli
                                                                                                                      Riley,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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                                                                                                                                                                                                                                                                                                                                        of
hypothetical C; Species: Me C; Date: 13-Se
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                                              H64489
                                                            RESULT
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Biosci. Biotechnol. Biochem. 60, 3
A;Title: Isolation and characteria
A;Reference number
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Description: probably a transcription regulator C;Superfamily: transcription regulator GntR C;Keywords: DNA binding; transcription regulation
                                 δ
                                                                                                                                                                                                                                                             C; Superfamily: alcohol dehydrog
C; Keywords: alcohol metabolism;
                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: GB:S51211; NID:g261968; PIDN:AAB24546.1; PID:g261969
A;Experimental source: strain DSM1617
A;Note: sequence extracted from NCBI backbone (NCBIN:120928, NCBIP:120938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                       F;174-203/Region: beta-alpha-beta NAD nucleotide-binding fold F;11:213/Modified site: N6-methyllysine (Lys) (partial) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: A; Residues: 1-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Accession: A44245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: A44245; MUID: 93099126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Function:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Gene: farR; gp30
A;Map position: 16.8 min
                                                                                                                                                                     F;98,101,104,112/Binding site:
                                                                                                                                                                                                                                               F;23-337/Domain: long-chain alcohol
                                                                                                                                                                                                                                                                                                     A; Pathway: alcohol degradation
                                                                                                                                                                                                                                                                                                                     A; Description: catalyzes the oxidation of primary and secondary alcohols to aldehydes
                                                                                                                                                                                                                                                                                                                                                                                 A; Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Title: Thermostable NAD(+)-dependent alcohol dehydrogenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R; Ammendola, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Accession: A44245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Date: 10-Mar-1994 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Species:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        alcohol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A44245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Experimental source: RU1012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Residues: 1-40 <UTS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Accession: PC4138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: GB:AE000176; GB:U00096;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                               Superfamily: alcohol dehydrogenase; long-chain alcohol dehydrogenase homology Keywords: alcohol metabolism; heat-stable protein; homodimer; metalloprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cross-references: DDBJ:D64014
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Experimental
                                                                         Matches
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                                                                                                                                                                                       8,68,154/Binding site:
220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  dehydrogenase (EC es: Sulfolobus sol
                                                                                                                                                                                                                                                                                                                                                                                 part of this sequence was
                               1 VLNASVTSPLLE
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VINASMODPLAE
                                                                                                                                                                                                                                                                                                                                                               homodimer
                                                                     Similarity 58.7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 58. 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                        1-347 <AMM>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Raia, C.A.; Caruso, C.; Camardella, L.; D'Auria, 12514-12523, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA
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231
                                 12
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                                                                                           58.3%;
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                                                                                                                                                                     zinc,
                                                                                                                                                                                       catalytic
                                                                                                                                                                                                                                                                                                                                                                                                   NCBI backbone (NCBIN:120928, NCBIP:120938)
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                                                                                         Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                 confirmed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kaino, Y.;
9-315, 1996
                                                                                                                                                                                                                                             dehydrogenase homology <LADH>
                                                                       ed. No. 36;
Mismatches
                                                                                                                                                                     noncatalytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sulfolobus solfataricus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                             Length 347;
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                                                                                                                                                                                       Cys) #status
                                                                                                                                                                                                                                                                   homodimer; metalloprotein;
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                                                                                                                                                                                                                                                                                                                                                                                 sequencing
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                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        from Sulfolobus solfatar
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protein MJ1521

Methanococcus jannaschii

revision

13-Sep-1996 #text\_change 13-Sep-1998

H64489; White,

Glodek, A.

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A;Title: Complete genome sequence of the methanogenic archaeon,
A;Reference number: A64300; MUID:96337999
A;Accession: H64489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rson, J.D.; Sadow, P.W.; Hanna, M.C.; Science 273, 1058-1073, 1996
                                                                                                       R;Fraser, C.M.; Norris, S.J.; Weinstock, G.M.; White, O.; Sutton, G.G.; Dodson, rson, J.; Khalak, H.; Richardson, D.; Howell, J.K.; Chidambaram, M.; Utterback, they, L.; Weidman, J.; Smith, H.O.; Venter, J.C.
                                                                                                                                                                   probable phosphoglucomutase - syphilis spirochete
C;Species: Treponema pallidum subsp. pallidum (syphilis spirochete)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 05-Nov-1999
C;Accession: C71327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ankyrin-like protein
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                                                                                            rson, J.; Khalak, H.; Richardson, they, L.; Weidman, J.; Smith, H.O
                                                                                                                                                                                                                                                                                                                                 В
                                                                                                                                                                                                                                                                                                                                                                  γQ
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                                        A;Title: Complete genome sequence of Treponema pallidum, the syphilis spirochete A;Reference number: A71250; MUID:98332770
                                                                               they, L.; Weidman, J.; Smi:
Science 281, 375-388, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            submitted to the Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R; Bevan, M.; Murphy, G.; Ridley, P.; Hudson, S.; Bancroft, I.; Mewes, H.W.; Rudd,
A; Status: preliminary; nucleic acid sequence not shown; translation not shown
                                      A; Reference number: A71250;
                                                                                                                                                                                                                                                                     RESULT
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                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 88.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Reference number: Z24493; Accession: T48596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Accession:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Molecule type: DNA; Residues: 1-422 <BUL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Status: preliminary; nucleic acid sequence not shown;
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Molecule type: DNA
Residues: 1-591 <BEV>
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                                                                                                                                                              Accession: C71
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Introns: 32/3
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Experimental source: cultivar
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78/3
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58.3%;
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 35;
Pred. No.
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45;
                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
67;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 591;
                                                                                                                                               O.; Sutton, G.G.; Dodson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Methanococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-Apr-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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McDd
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                       Q
                                                                                   Query Match
Best Local
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Local Similarity nes 6; Conserv

Conservative

63.6%;

Score 35; DB Pred. No. 91; 2; Mismatches

1;

Indels

0;

Gaps

0

DВ 1;

1 VLNASVTSP

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A;Molecule type: DNA
A;Residues: 1-632 <COL>
A;Cross-references: GB:AE001219; GB:AE000520; NID:g3322693; PIDN:AAC65401.1; PID:g332
A;Experimental source: strain Nichols
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Date: 10-Sep-1999 #sequence
C; Accession: S61642; S20179;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Gene:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Vambutas, A.; Ackerman, S.H.; Tzagoloff, A.
Eur. J. Blochem. 201, 643-652, 1991
A;Title: Mitochondrial translational-initiation and elongation factors in Saccharomyc
A;Reference number: S20177; MUID:92037620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               translation elongation factor EF-G, mitochondrial - N;Alternate names: protein L2195; protein YLR069c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                S61642
                                                                                                                                     C;Keywords: GTP binding; mitochondrion; P-loop; protein bic F;71-205/Domain: translation elongation factor Tu homology
                                                                                                                                                                                                                                                                                                 A;Cross-references: EMBL:Z73241; NID:g1360421; PIDN:CAA97626.1; PID:g1360422; GSPDB:G
A;Experimental source: strain S288C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           submitted to the Protein Sequence Database, May 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-65, 'I',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  submitted to the EMBL Data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Urrestarazu,
                                                                    F;327-329/Region:
                                                                                                                                                                                                                                                        A;Gene: SGD:MEF1; MIPS:YLR069c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-761 <AND>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Reference number: S64872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      a;Residues: 1-65,'I',67-232,'V',234-477,'S',479-628,'P',630-761
a;Cross-references: EMBL:X58378; NID:93917; PIDN:CAA41267.1; PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:X94607; NID:g1181264; PIDN:CAA64315.1; PID:g1181289
R;Yambutas, A.; Ackerman, S.H.; Tzagoloff, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-761 <URR>
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                                                                                                                                                                                   a; Map position: 12R
C; Superfamily: translation elongation factor G;
                                                                                                                                                                                                                                  A;Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Experimental source:
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                        W; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Species: Saccharomyces cerevisiae
                                                                                  77-84/Region: nucleotide-binding motif A (P-loop) 202-205/Region: GTP-binding NKXD motif
                                                                                                                                                                                                                                                                                  Genetics:
                                                                                                                                                                                                                                                                                                                                                                     Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cross-references: EMBL: Z73241;
                                                                                                                                                                                                                                                                                                                                                                                                  Accession: S64901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          182 LNASILEPLL 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ν
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                                                                                                                                                                                                                                                                                                                                                                                                                                            to the Protein
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7; Conserv
                                                                                                                                                                                                                                                                                                                                                  1-761 <POH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S2017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Urrestarazu, L.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1999 #sequence_revision 10-Sep-1999 #text_change 21-Jul-2000
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                                                                      GTP-binding SAK/L motif
                                                                                                                                                                                                                                                                                                                                                                                                                      otein Sequence Database,
S64899
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                                                                                                                                                                                                                                  SGD:S0004059; MIPS:YLR069c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            strain
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          41; NID:g1360421; PIDN:CAA97626.1; PID:g1360422; GSPDB:G
S288C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ŗ
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Pred. No. 73;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    December 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                   translation elongation factor Tu homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            yeast (Saccharomyces cerevisiae)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 632
Length 761
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PID: 93918
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A; Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Smith, H.O.; Woese, C.R.; Venter, J.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: C69304
R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson R;Klenk, H.P.; Clayton, R.A.; Tomb, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probable enoy1-CoA hydratase (EC 4.2.1.17) fad-1 - Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Date: 28-May_1999 #sequence_revision 28-May-1999 #text_change 18-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
6; Conserv
C; Species: Mus muscu
C; Date: 02-Jul-1996
                                                                     RESULT
148471
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                       Fc gamma (IgG) receptor high affinity - mouse C; Species: Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PHO85 protein
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N;Reference number: S61996
N;Accession: S62011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;Alternate names:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Title: The complete genome sequence of the hyperthermophilic, Reference number: A69250; MUID:98049343
Accession: C69304
                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1110-1126/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene: PHO85; MIPS:YPL110c Map position: 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schlenstedt,
                                                                                                                                                                                                                                                                                                                                                                Cross-references: GB:AE001074; GB:AE000782; NID:g2689397; PIDN:AAB90798.1; PID:g265019 Superfamily: naphthoate synthase; encyl-CoA hydratase homology Keywords: carbon-oxygen lyase; fatty acid beta-oxidation; hydro-lyase 25-175/Domain: encyl-CoA hydratase homology <ECH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Superfamily: unassigned ankyrin repeat proteins; ankyrin
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Residues: 1-1223 <SCH>
Cross-references: EMBL:U43503; NID:g1163087; PIDN:AAB68251.1;
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Residues: 1-256 <KLE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Keywords: transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genetics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            i77-593/Domain: transmembrane #status predicted <TM1>
i110-1126/Domain: transmembrane #status predicted <TM2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tatus: nucleic acid sequence not shown; translation not shown
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                                                                                                                                                                                                                                                         Conservative
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#sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           #sequence_revision 19-Apr-1996 #text_change 05-Nov-1999
                                                                                                                                                                                                                                                                            61.8%;
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54.5%;
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Pred. No.
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Pred: No. 40;
1; Mismatches
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                                                                                                                                                                                                                                                                                               Length 256;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            J.F.; McDonald,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sulfate-reducing archae
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Best Local Similarity
Matches 8; Conserve
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                                                                                                                                                                                                                                                                                                         A;Title: Linkage on chromosome 3 of autoimmune diabetes and defective Fc receptor for A;Reference number: I48471; MUID:93242399
A;Accession: I48471
                                                                                                                                                                                                                                                                                                                                                                   C; Accession: I48471
R; Prins, J.B.; Todd, Science 260, 695-698,
                                                                                                                                                                                                                                                      A; Residues:
                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                         A;Status: preliminary; translated from GB/EMBL/DDB:
207 VLRASVSSPFPE 218
                                      1 VLNASVTSPLLE 12
                                                                                                                                                                                                                                                    1-336 <RES>
                                                                                                                                                                                                                                                                                                                                                                                       Todd, J.A.;
                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                       1993
                                                                                            61.8%;
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                                                                                              Score 34; DB Pred. No. 55;
                                                                            Mismatches
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                                                                                                                Length 336
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Search completed: June 4, 2001, 12:15:38 Job time: 271 sec

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Result
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"Nuclectide sequence of three cDNAs for receptor (FCRI)."; Nucleic Acids Res. 16:11824-11824(1988). [2] [2] SEQUENCE FROM N.A. MEDLINE-89100284; PubMed=2911749; Allen J.M., Seed B.; "Isolation and expression of functional complementary DNAs."; Science 243:378-381(1989)!- FUNCTION: BINDS TO THE FC REGION OF AFTINITY RECEPTOR!- SUBCELLULAR LOCATION: TYPE I MEMBRAN PRODUCTS: 2 ISOFORMS; A PRODUCED BY ALTERNATIVE SPLICING!- TISSUE SPECIFICITY: MONOCYTE/MACROPH -!- SIMILARITY: CONTAINS 3 IMMUNOGLOBULI!- DATABASE: NAME-PROW; NOTE-CD guide C WMW-"http://www.ncbi.nlm.nih.gov/pro This SWISS-PROT entry is copyright. It i	RESU POC OCC OCC OCC OCC OCC OCC OCC OCC OCC	33 34 32 58.2 35 32 58.2 36 32 58.2 37 32 58.2 38 32 58.2 40 32 58.2 41 32 58.2 42 32 58.2 43 32 58.2 44 32 58.2 45 32 58.2 47 32 58.2 48 32 58.2 49 32 58.2 40 32 58.2 41 32 58.2 42 32 58.2 43 32 58.2 45 32 58.2 46 32 58.2 47 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
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J. Bacteriol. 178:301-305(1996).

I CATALYTIC ACTIVITY: ALCOHOL + NAD(+) = ALDEHYDE OR K-1- COFACTOR: BINDS FOUR ZINC IONS PER DIMER.

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Pred. No. 0.0017;
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RI tissue-specific expression.";

RL FEBS Lett. 416:30-34(1997).

RL FEBS LETT. THE SUBCLET LIS TO SUBSTRATES CONTAINING PROMOTES DOCKING OF IMPORT COMPONENTS SEPARATE AND IMPORTIVE ALPHA AND -BETA ARE RECYPOPLASMIC SIDE OF THE NPC, THE COMPONENTS SEPARATE AND IMPORTIVE ALPHA AND -BETA ARE RECYPOPLASMIC.

CC THREE COMPONENTS SEPARATE AND IMPORTIVE ALPHA AND -BETA ARE RECYPOPLASMIC.

CC -ITSSUE SPECIFICITY: ONLY SIGHTLY DETECTED IN EHRLICH ASCITES THE SEPARATE AND SKELETAL MUSCLE, CLEARLY DETECTED IN CHUNCOLUMN SALUENT SEPARATE, AND LUNG. HIGH EXPRESSION IN TESTIS.

CC -IUNDROLUMNIS AND SKELETAL MUSCLE, CLEARLY DETECTED IN CHUNCOLUMN AND SEPARATE AND A SHORT
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
IMPORTIN ALPHA-6 SUBUNIT (KARYOPHERIN ALPHA-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOUSE
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HSSP; P00325;
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                                                                                                                                                                                                                                                                                                                                           EDLINE=98034160; ruumeu->>>>--. Yoneda Y.; suji L., Takumi T., Imamoto N., Yoneda Y.; Identification of novel homologues of mouse importin alpha, the Identification of novel homologues of mouse importin alpha, the
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                                             SIMILARITY: BELONGS TO THE IMPORTIN SIMILARITY: CONTAINS 8 ARM REPEATS.
                                                                                                     HYDROPHOBIC CENTRAL REGION COMPOSED OF 8 REPEATS, AND A HYDROPHILIC C-TERMINUS. THE N-TERMINAL HYDROPHILIC REGIC THE IMPORTIN BETA BINDING DOMAIN (IBB DOMAIN), WHICH IS
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Sciurognathi; Muridae
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European Bioinformatics Institute.

There are no restrictions on ng as its content is in no

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STRAIN-K12 / MG1655;
MEDLINE-97426617; PubMed-9278503;
Blattner F.R., Plunkett G. III, Bloch C.A.,
Riley M., Collado-Vides J., Glasner J.D., R
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01-NOV-1997 (Rel.
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01-JAN-1990 (Rel. 13, Last seno
01-NOV-1997 (Rel. 35, Last anno
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                                          SEQUENCE OF 1-40 FROM N.A. MEDLINE=97076625; PubMed=9
                                                                                                                                                          Gregor J., Davis N.W., Kirkpatrick H.A.,
                                                                                                                                                                                                                                                                                                                                                                       Buck D.,
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Bacteria; Proteobacteria;
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PROSITE; PS50176;
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                                                                                                complete genome sequence of 
nce 277:1453-1474(1997).
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Last annotation update)
REGULATOR (P30 PROTEIN)
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Kaino Y., Tanabe H., Noda M.;
n of the heat-responsive genes
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                                                                                                                  Escherichia coli K-12.";
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                                                                                                                                                          Goeden M.A.,
                                                                                                                                                                          Rode C.K., Mayhew G.F.,
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                                                                                                                                                                                               Perna N.T.,
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                                                                                                                                                                                                                                                                                                                               of the succinyl-CoA sequence of a gene
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                                                                                                                                                          D.J.,
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Best Local
                                             ADH1_SULSO
P39462; O740
O1-FEB-1995
                                                                                        SULSO
01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
NAD-DEPENDENT ALCOHOL DEHYDROGENASE (EC 1.1
                                                                                                                                                                                                                                                                                           Interpro; IPR000524; -
Pfam; PF00392; gntR; -
Pfam; PF00392; HTHGNTR.
PRINTS; PR00035; HTHGNTR.
PROSTIE; PS00043; HTH_GNTR.FAMILY; 1.
Transcription regulation; DNA-binding.
DNA_BIND 32 51 H-T-H MOTIF (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ikemoto K., Inada H., Baba T., Fujita K., Hayashi K., Honjo A., Ikemoto K., Inada T., Itoh T., Kajihara M., Kanai K., Kashimoto K Kimura S., Kitagawa M., Makino K., Masuda S., Miki T., Mizobuchi Mori H., Motomura K., Nakamura Y., Nashimoto H., Nishio Y., Saito Sampei G., Seki Y., Tagami H., Takemoto K., Wada C., Yamamoto Y., Yano M., Horiuchi T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEBS Lett. 356:183-187(1994)

1: FUNCTION: POSSIBLE TRANSCRIPTIONAL REGULATOR OF THE SUCCINYL-COA-
SYNTHEFFASE OPERON. BINDS SPECIFICALLY TO THE FARR PROMOTER AND
EXHIBITS WEAK BINDING TO THE CITRATE SYNTHASE AND LIPOAMIDE
DEHYDROGENASE PROMOTERS. BINDING AT THE FARR PROMOTER IS ABOLISHED
BY LONG-CHAIN FATTY ACIDS AND THEIR COA THIOESTERS. NEGATIVELY
BY LONG-CHAIN FATTY OWN SYNTHESIS IN RESPONSE TO LONG-CHAIN FATTY
                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=95104410; PubMed=7805834; Quail M.A., Dempsey C.E., Guest J.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "A 718-kb DNA sequence of the corresponding to the 12.7-28.0
                                                                                                                                                                                                                                                                                                                                                                                     EMBL; D90711; BAA35396.1; PIR; S04645; S04645. EcoGene; EG11109; farR.
                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; D64014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; X15790; CAA33791.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Identification of
Escherichia coli.";
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                                                                                                                                                                 1 YLNASVTSPLLE 12 :|| | 1 | | | |
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SIMILARITY: BELONGS TO THE GNTR FAMILY OF TRANSCRIPTIONAL
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                                                                074076;
                                                                                                                                                                                                          Similarity 58. 7; Conservative
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                                                                                                                                                                                                                                                                                    240 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                 -; NOT_ANNOTATED_CDS
                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PubMed=8905232;
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                                                                                                                                                                                                                        Score 35;
Pred. No.
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                                                                             PRT.;
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                                                                                                                                                                                                                          DВ
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on the lin
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L outstation -
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on update) (EC 1.1.1.

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SOT TITT TEXTOR REPRESENTATION OF THE PRESENTATION OF THE PRESENTA
            SSETTTAL
                                                                                                                                                                  GRHR_CLAGA
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Best Local S
Matches 7
GRHR_CLAGA STANDARD,

GRHR_CLAGA STANDARD,

042329;

15-JUL-1998 (Rel. 36, Created)

15-JUL-1998 (Rel. 36, Last sequence update)

15-JUL-1998 (Rel. 36, Last annotation update)

GONADOTROPIN-RELEASING HORMONE RECEPTOR (GNRH
GONADOTROPINS (Sharptooth catfish) (Afri
                                                                                                                                                                                                                                                                                                                                                                                                                      METAL
METAL
SEQUENCE
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relationship to other alcohol dehydrogenases.";
Biochemistry 31:12514-12523(1992);
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METAL
METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=DSM 1617 / P2;
Aravalli R.N.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sulfolobus solfataricus Archaea; Crenarchaeota; NCBI_TaxID=2287;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oxidoreductase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; S51211; AZ
EMBL; AJ010590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          between the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ammendola S., Raia C.
de Rosa M., Rossi M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=93099126; PubMed=1463738;
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STRAIN-DSM 1617 / P
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                                                                                                                                                                                                                                               220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mitted (AUG-1998) to the EMBL/GenBank/DDBJ databases CATALYTIC ACTUTY: ALCOHOL + NAD(+) = ALDEHYDE OR COFACTOR: BINDS FOUR ZINC IONS DER DIMER. SUBUNIT: HOMODIMER AND HOMOTETRAMER.
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58.3%;
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METHYLATION
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          (African catfish)
Vertebrata; Eutel
                                                (GNRH-R).
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            Euteleostomi;
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"Distinct efficacies for two endogenous
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FUNCTION: THIS RECEPTOR MEDIATES ITS ACTION BY ASSOCIATION WITH G PROTEINS THAT ACTIVATE A PHOSPHATIDYLINOSITOL-CALCIUM SECOND
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Sutton G.G., Blake J.A., Fitzgerald L.M., Clayton R.A., Gocayne J.D.,
Kerlavage A.R., Dougherty B.A., Tomb J.-F., Adams M.D., Reich C.I.,
Overbeek R., Kirkness E.F., Weinstock K.G., Merrick J.M., Glodek A.,
Scott J.L., Geoghagen N.S.M., Weidman J.F., Finhrmann J.L., Nguyen D.,
Utterback T.R., Kelley J.M., Peterson J.D., Sadow P.W., Hanna M.C.,
Cotton M.D., Roberts K.M., Hurst M.A., Kaine B.P., Borodovsky M.,
Klenk H.-P., Fraser C.M., Smith H.O., Woese C.R., Venter J.C.,
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EFG1\_YEAST
ID EFG1\_Y
AC P25039
DT 01-MAY

EFG1\_YEAST S P25039; Q99360; 01-MAY-1992 (Rel

(Rel. 22, Created)

STANDARD;

PRT;

761 AA

RESULT

Вb

205 VLNCEILPPLLE

216

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Q.
                 Query Match
Best Local Similarity
Thes 7; Conserv
                                                                                              DOMAIN
SEQUENCE
                                                                                                                                                                                                                                                                                                           Pfam;
                                                                                                                                                     REPEAT
REPEAT
                                                                                                                                                                                                            REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=98055463; PubMed=9395085;
                                                                                                                                                                                REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                        MIM; 604545;
                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF005361;
                                                                                                                                                                                                                                                                                                                                                                                              or send
                                                                                                                                                                                                                                                                                                                                                                                                              entities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute.
use by non-profit institutions as lo
modified and this statement is not remo
                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the expression pattern of the importing alpha protein family FEBS Lett. 417:104-108(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Roehler M., Ansieau S., Prehn S., Leutz A., Haller H., Hartmann i "Cloning of two novel human importin-alpha subunits and analysis the expression pattern of the importin-alpha protein family.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Mammalia; Eutheria;
                                                                                                                                         REPEAT
                                                                                                                                                                                                                                                                                 PROSITE;
                                                                                                                            DOMAIN
                                                                                                                                                                                                                                        REPEAT
                                                                                                                                                                                                                                                                                                                             InterPro;
                                                                                                                                                                                                                                                                                                                                        InterPro; IPR000225;
                                                                                                                                                                                                                                                        DOMAIN
 1 VLNASVTSPLLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: BINDS SPECIFICALLY AND DIRECTLY TO SUBSTRATES CONTAINING EITHER A SIMPLE OR BIPARTITE NLS MOTHE, PROMOTES DOCKING OF IMPORT SUBSTRATES TO THE NUCLEAR PORE COMPLEX (NPC). THE COMPLEX IS SUBSEQUENTLY TRANSLOCATED THROUGH THE PORE BY AN ENERGY REQUIRING, RAN-DEPENDENT MECHANISM. AT THE NUCLEOPLASMIC SIDE OF THE NPC, THE RAN-DEPENDENT MECHANISM. AT THE NUCLEOPLASMIC SIDE OF THE NPC, THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY:
SIMILARITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION: CYTOPLASMIC. TISSUE SPECIFICITY: TESTIS.

DOMAIN: CONSISTS OF AN N-TERMINAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DOMAIN: CONSISTS OF AN N-TERMINAL HYDROPHILIC REGION, A HYDROPHOBIC CENTRAL REGION COMPOSED OF 8 REPEATS, AND A SHORT HYDROPHILIC C-TERMINUS. THE N-TERMINAL HYDROPHILIC REGION CONTAINS THE IMPORTIN BETA BINDING DOMAIN (IBB DOMAIN), WHICH IS SUFFICIENT FOR BINDING IMPORTIN BETA AND ESSENTIAL FOR NUCLEAR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SUBUNIT: FORMS A COMPLEX WITH IMPORTIN BETA-1 SUBUNIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   THREE COMPONENTS SEPARATE AND IMPORTIN-ALPHA AND -BETA ARE EXPORTED FROM THE NUCLEUS TO THE CYTOPLASM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IMPORT
                                                                                                                                                                                                                                                                                              PF01749;
                                                                                                                                                                                                                                                                                                           PF00514; Armadillo_seg;
                                                                                                                                                                                                                                                                                                                                                                                                an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                  PS50176;
                                                                                                                                                                                                                                                                                                                             IPR002652;
                                                                                             411
25
454
536 AA;
                                                                                                                                                                                                                                                                     Protein
                             Conservative
                                                                                                                                                                                                                                                                                                 IBB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BELONGS TO THE IMPORTIN ALPHA FAMILY. CONTAINS 8 ARM REPEATS.
                                                                                                                                                                                                                                                                                                                                                                    AAC51868.1;
  12
                                                                                                                                                                                                                                                                                 ARM_REPEAT;
                                                                                                                                                                                                                                   159
                                                                                                            410
453
28
536
                                                                                                                                                                     288
328
370
                                                                                                                                                                                                                                                                  transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chordata;
Primates;
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                                                                                                60266
                                          58.3%;
                                                                                                MW;
                                                                                                                                      IBB
ARM
ARM
ARM
ARM 4
ARM 5
ARM 6
ARM 7
ARM 7
                             1.
                                        Score 35; DB
Pred. No. 26;
                                                                                       POLY-ARG.
ASP/GLU-RICH (ACIDIC).
, 4E880B1FAD2A420E CRC64;
                                                                                                                                                                                                                                                                                                             8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                    Repeat
                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                             (See http://www.isb-sib.ch/announce/
                                                       1:
                                                       Length.536;
                                                                                                                                                                                                                                                                                                                                                                                                                            and
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                             Gaps
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ELONGATION FACTOR 15-JUL-1998 (Rel. 15-JUL-1999 (Rel.

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RESULT 10
FTSZ_AQUAE
ID FTSZ_AQUAE
                                                          日
                                                                                                                                                                    RT RXX
                                                                                                                    Query Match
Best Local
                                                                                                          Matches
                                                                                                                                                                              NP_BIND
NP_BIND
CONFLICT
CONFLICT
CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Andre B., Urrestarazu L.A.;
Andre B., Urrestarazu L.A.;
Submitted (MAY-1996) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: THIS PROTEIN PROMOTES THE GTP-DEPENDENT TRANSLOCATION
OF THE NASCENT PROTEIN CHAIN FROM THE A-SITE TO THE P-SITE OF TH
                                                                                                                                                                                                                                                                                                    PRINTS; PRO0315; ELONGATNECT.
PROSITE; PS00301; EFACTOR_GTP; 1
Elongation factor; Protein biosy
                                                                                                                                                                                                                                                                                                                                        InterPro; IPR000640; -.
InterPro; IPR000795; -.
Pfam; PF00679; EFG_C; 1.
Pfam; PF00009; GTP_EFTU;
                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; x58378; CAA41267.1;
EMBL; x94607; CAA64315.1;
EMBL; Z73241; CAA97626.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae (Baker's yeast).
                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                   CHAIN
                                                                                                                                                                                                                                                                              TRANSIT
                                                                                                                                                                                                                                                                                       Transit peptide;
                                                                                                                                                                                                                                                                                                                                                                                      HSSP; Pl3551; 1ELO.
SGD; S0004059; MEF1.
                                                                                                                                                                                                                                                                                                                                                                                                              PIR; S17025;
PIR; S20179;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomyces cerevisiae.";
Eur. J. Biochem. 201:643-652(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=92037620; Pub
Vambutas A., Ackerman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomycetales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Fungi; Ascomycota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEF1 OR YLR069C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-92037620; PubMed=1935960;
Vambutas A., Ackerman S.H., Tzagoloff A.;
"Mitochondrial translational-initiation and elongation factors in
                                                           899
                                                                                                          Local Similarity hes 6; Conser
                                                                                 1 VLNASVTSP 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: MITOCHONDRIAL SIMILARITY: BELONGS TO THE GTP-BINDI EF-G/EF-2 SUBFAMILY.
                                                           IMNVSVTSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and this statement is not removed requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    email to license@isb-sib.ch)
                                                                                                                                                                                           148
202
66
233
478
                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              S17025.
                                                                                                                                                                    AA;
                                                          676
                                                                                                                                                                                                                                                                                                                                        GTP_EFTU; 1
 STANDARD;
                                                                                                                                                                                                                                                                                        GTP-binding.
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38,
1
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84
152
205
205
233
478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                    Protein biosynthesis; Mitochondrion;
                                                                                                                                                                    629
84573
                                                                                                                    63.6%;
66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Last sequence update)
Last annotation update)
L MITOCHONDRIAL PRECURSOR (MEF-G-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TO THE GTP-BINDING ELONGATION FACTOR FAMILY
                                                                                                                                                                    MW;
                                                                                                                                                                                         GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
R -> I (IN REF. 1).
P -> S (IN REF. 1).
                                                                                                                  Score 35;
Pred. No.
                                                                                                                                                                                                                                                                             MITOCHONDRION (POTENTIAL)
  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Saccharomycotina; Saccharomycetes;
                                                                                                                                                                    -> P (IN REF. 1) 3E2C534509B09103
                                                                                                       Mismatches
 367 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (See http://www.isb-sib.ch/announce/
                                                                                                                     39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  There
                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as its content
                                                                                                        1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  are no
                                                                                                                               Length 761;
                                                                                                                                                                    CRC64;
                                                                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL
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RESULT
FCG1_MC
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                                                                                                                                   MOUSE
MOUSE STANDARD; PRT; 404 AA.

FCGL_MOUSE STAN
             Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRINTS; PR00423; CELLDVISFTSZ.

PROSITE; PS01134; FTSZ_1; 1.

PROSITE; PS01135; FTSZ_2; 1.

Cell division; Septation; GTP-binding.

Cell division; Septation; GTP (POTENTIAL).

NP_BIND 100 GTP (POTENTIAL).

SEQUENCE 367 AA; 40161 MW; ODECBA6A865348A9 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nature
-!- FUN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aquifex aeolicus.
Bacteria; Aquificales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAY-2000 (Rel. 39, 30-MAY-2000 (Rel. 39, 30-MAY-2000 (Rel. 39,
                                                                                              Mus musculus (Mouse).
                                                                                                                             FCGR1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entities requires a license agreement (or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Deckert G., Warren P.V., Gaasterland T., Young W.G., Lenox A.L., Graham D.E., Overbeek R., Snead M.A., Keller M., Aujay M., Huber Feldman R.A., Short J.M., Olson G.J., Swanson R.V.; "The complete genome of the hyperthermophilic bacterium Aquifex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FTSZ OR AQ_525.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; AE000694; AAC06771.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=98196666; PubMed=9537320;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN-VF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=63363;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00091; tubulin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR000158; InterPro; IPR003008;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A
                                                                                                                             ) (FCRI) (IG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 SVTSPLLE 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY: BELONGS TO THE FTSZ FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BEGIN BINDS TO AND HYDROLYZES GTP (BY SIMILARITY).
SUBUNIT: AGGREGATE TO FORM A RING-LASSEMBLE AT THE INNER SURFACE
OF THE CYTOPLASMIC ASSEMBLE AT THE INNER SURFACE
OF THE CYTOPLASMIC MEMBRANE (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: THIS PROTEIN IS ESSENTIAL TO THE CELL-DIVISION PROCESS. ITS SEEMS TO ASSEMBLE INTO A DYNAMIC RING ON THE INNER SUFFACE OF THE CYTOPLASMIC MEMBRANE AT THE PLACE WHERE DIVISION WILL OCCUR, AND THE FORMATION OF THE RING IS THE SIGNAL FOR SEPTATION TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AVTSPLLE 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DIVISION PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            392:353-358(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61.8%;
87.5%;
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Last annotation update)
FTSZ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 34;
Pred. No.
                                            Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (See http://www.isb-sib.ch/announce/
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27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 367;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        restrictions
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-!- TISSUE SPECIFICITY: MACROPHAGE-SPECIFIC.
-!- SIMILARITY: CONTAINS 3 IMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
-!- CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-10 IS THE INITIATOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=92166399; PubMed=1531670;
Osman N., Kozak C.A., McKenzie I.F., Hogarth P.M.;
"Structure and mapping of the gene encoding mouse
gamma RI and chromosomal location of the human Fc
J. Immunol. 148:1570-1575(1992).
                                                                                                       01-NOV-1995
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01-OCT-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration
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Yarrowia lipolytica (Candida lipolytica)
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A46480; A46480.
; P12319; IALT.
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Osman N., Tate B., McKenzie I.F.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    144:371-378(1990).
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IG-LIKE C2-TYPE DOMAIN (TRUNCATED).
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IG-LIKE C2-TYPE DOMAIN
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POTENTIAL.
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of the mouse high affinity Fc
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GLE1_YEAST
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Interpro; IPR00100...

Pfam; PF00649; Copper-tls.,

Pfam; PF00649; CopperFIST.

PRINTS; PR00617; CopperFIST_1; 1.

PROSITE; PS0073; COPPER_FIST_2; 1.

PROSITE; PS0073; COPPER_FIST_2; 1.

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SEQUENCE
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                                                                                                                                                                                                 Murphy R., Wente S.R.; "An RNA-export mediator with an essential nuclear export signal."; Nature 383:357-360(1996).
                                                                                                                                                                                                                                                         SEQUENCE FROM N.A., MEDLINE=96404361; P
                                                                                                                                                                                                                                                                                                                                                            Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
                                                                                                                                                                                                                                                                                                                                                                                                    GLE1 OR BRR3 OR YDL207W OR D1049
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15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or send an email to license@isb-sib.ch).
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                                                                                                  Submitted
                                                                                                                   Bahr A.,
                                                                                                                                      STRAIN-S288C
                                                                                                                                                                                                                                                                                                                                           Saccharomycetales;
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              AIN-5288C / FY1679;

X A., Moeller-Rieker S., Hankeln T., Kraemer C., Schmidt E X A., Moeller-Rieker S., Hankeln T., Kraemer C., Schmidt E mitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.

FUNCTION: RNA EXPORT FACTOR. THE MECHANISM MAY INVOLVE TH TARGETING OF RNA/RNP MOLECULES TO OR THROUGH THE NUCLEAR COMPLEX BY MEANS OF RIP1 AND NUP100 INTERACTIONS.

SUBCELLULAR LOCATION: NUCLEAR. LOCALIZED PREDOMINANTLY AT
NUCLEAR PORE COMPLEXES
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                                                                                                                                                                                                                                                         PubMed=8848052;
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                                                                                                                                                                                                                                                                                                                                           Saccharomycetaceae;
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Pred. No.
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GBA4_USTMA
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Best Local
NP_BIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "G proteins in Ustilago maydis: transmission of multiple signals?",
EMBO J. 16:1934-1942(1997).

-1- FUNCTION: GUANRINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE
INVOLVED AS MODULATORS OR TRANSDUCERS IN VARIOUS TRANSMEMBRANE
SIGNALING SYSTEMS.

-1- SUBUNIT: G PROTEINS ARE COMPOSED OF 3 UNITS (ALPHA, BETA & GAMM
THE ALPHA CHAIN CONTAINS THE GUANINE NUCLEOTIDE BINDING SITE.

-1- SIMILARITY: BELONGS TO THE G-ALPHA FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SGD;
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                                                                                      Pfam; PF00503; G-alpha; 1. PRINTS; PR00318; GPROTEINA.
                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restricted the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement of the statement 
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01-OCT-2000 (Rel.
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P87035;
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                                                         GTP-binding;
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Ustilaginomycetidae;
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                                                                                                                                                                                                                                                                      and this statement is not removed. Usage by and for commercial requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                             non-profit institutions as long
                                                                                                                                                   IPR001019; -.
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                                                         Transducer;
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i; Basidiomycota;
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                                                            Multigene family
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; C9C6B1513AF95711 CRC64;
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L->A: PARTIAL LOSS OF ACTIVITY.
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(BY SIMILARITY).
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Best Local
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Q10197; P78749;
01-OCT-1996 (Rel. 34, Crea
15-JUL-1999 (Rel. 38, Last
15-JUL-1999 (Rel. 38, Last
TUBUFLIN-FOLDING COFACTOR D
ALPI OR SPBC11C11.04C.
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SEQUENCE
                                                                                                                                                                                                                                    use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                           Wood V., Rajandream M.A., Barrell B.G., Beck A., Rei
Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databa-
i- FUNCTION: HAS A FUNCTION IN THE FOLDING OF BETA-
A VITAL ROLE IN MICROTUBULE-DEPENDENT PROCESSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SCHPO
                                                       CARBOHYD
                                                                                                                                                                              EMBL;
                                                                                                                                                                                                                                                                                                                                                             STRAIN-972
Wood V., R
                                                                                                                                                                                                                                                                                                                                                                                                     Hirata D., Masuda H., Eddison M., Toda T.; "Essential role of tubulin-folding cofactor D in microtubule and its association with microtubules in fission yeast."; EMBO J. 17:658-666(1998).
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MEDLINE=98119809; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetes;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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J. Clin. Invest. 90:2102-2109(1992).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      J. Exp. Med.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gamma receptor gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Definition of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (DEC-1992) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Benech P.D., Sastry K.N., Iyer R.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-FEB-1997
                                                                                                                                                                                                  36.6 KDA PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NTERPRO; IPRO03006; -. 

PAM; PF00047; ig; 3. 

RODOM; PD002534; -; 1.
                                                                                                                                                                                                                                                                                                                                                       194
                 EQUENCE OF 1-7 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                   VLNASVTSPLLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P12319;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S45708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S4570
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       M915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S45705;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S45704;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             M9155
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         M91550;
                                                                                                                                                                                                                                                                                                                                                                                                                 l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A.B.
                                            (MAR-1995)
                                                                                                                                                                                                                (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                                              (TrEMBLrel.
                                                                                                                                                                                                                                                                            PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    176:1115-1123(1992)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA58414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1ALT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD13842
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA58414.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA58414.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                interferon gamma-response elements in a novel human Fc gene (Fc gamma RIb) and characterization of the gene
                                                                          deduced amino aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
                                                                                                                                                                                                                                                                                                                                                                                  12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42881 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                             ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02,
15,
                                               the
                                                                                                                                                                    Reoviridae; unclassified Reoviridae
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JOINED.
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                                                             of
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                                             EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                Last
Last
                                                                                                                                                                                                                                             Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 55; DB 4; Pred. No. 0.0094;
                                                             the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A84D464C70DD0F91 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                sequence up
                                                            sequences of Nilaparvata lugens
genus Fijivirus.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eichbaum
                                                                                                                                                                                                                                                                            1199
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                                                                                                                                                                                                                                  update)
                                                                                                                                                                                                                úpdate)
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                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 375;
                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Raveh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Raveh
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P7911
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Best Local S
Matches 8
       REPEAT
REPEAT
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REPEAT
REPEAT
                                                                                                              DOMAIN
                                                                                                                                              PROPEP
CHAIN
                                                                                                                                                                                                       PFAM; PF00560; LRR; 3.
PFAM; PF01462; LRRNT; 1.
Proteoglycan; Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Characterizatto... _ "The repeat core protein."; rich repeat core protein."; rich repeat core protein."; rich repeat core protein."; aND _ _ _ _ _ _ _ A ROLE IN BONE FORMATION, AND _ _ _ _ _ _ _ OF CARTILAGE 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-1997 (Tremblrel. 03,
01-MAY-1997 (Tremblrel. 03,
01-OCT-2000 (Tremblrel. 15,
SMALL CHONDROITIN/DERMATAN S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P79119;
P79119;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            putative member of the genus Fi
J. Gen. Virol. 77:139-146(1996)
EMBL; D49694; BAA08543 1; ...
                                                                                                                                                                                                                                                                                  MANY PROTEINS. NUMBER IN THIS -i- SIMILARITY: IN THE C-TERMINAL EMBL; U77127; AAB68397.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A., PARTIAL SEQUENCE, TISSUE=FETAL EPIPHYSEAL CARTILAGE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nakashima N., Koizumi M., Watanabe H., Hiroaki N., "Complete nucleotide sequence of the Nilaparvata l putative member of the genus Fijivirus.";
                                                                                                                                                                            SIGNAL
                                                                                                                                                                                                                                                                                                                                                               <del>-</del>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=97373567; PubMed=9228042;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bovidae; Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=96145144; PubMed=8558122;
                                                                                                                                                                                                                                                                    INTERPRO; IPRO00372; -
                                                                                                                                                                                                                                                                                                                                  +
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PGLB) (EPIPHYCAN)
                                                                                                                                                                                              Connective
                                                                                                                                                                                                                                                       INTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                             ÷
                                                                                                                                                                                                                                                                                                                                                                                                                                          ÷
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Johnson H.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           753 VLNTLVTSPIVE
                                                                                                                                                                                                                                                                                                     (POTENTIAL).
SIMILARITY: THE REPEATED LEUCINE-RICH (LRR)
MANY PROTEINS. NUMBER IN THIS PROTEIN: 6.
                                                                                                                                                                                                                                                                                                                                                             PTM: A LONG AND A SHORT FORM; PRESENT IN APPROXIMATELY EQUIMOLAR AMOUNTS; MAY ARISE BY PROTEOLYSIS OR CLEAVAGE BY EXOPEPTIDASES. PTM: THE THREE GLYCOSAMINOGLYCAN CHAINS ARE DERMATAN SULFATE
                                                                                                                                                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: EXTRACELLULAR MATRIX (BY SIMILARITY). TISSUE SPECIFICITY: PREFERENTIALLY EXPRESSED IN THE ZONE OF FLATTENED CHONDROCYTES OF THE DEVELOPING LIMB CARTILAGE. DEVELOPMENTAL STAGE: EMBRYO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VLNASVTSPLLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               h 72.7%;
Similarity 66.7%;
8; Conservative
                                                                                                                                                                                                                                                       IPR001611; -.
                                                                                                                                                                                            tissue;
         152
167
191
191
211
237
258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
                                                                                                                                                 20
31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rosenberg L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           764
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                       166
190
210
236
257
288
                                                                                                                                                                                            Extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cetartiodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               136614 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 , Last sequence update)
, Last annotation update)
SULFATE PROTEOGLYCAN PRECURSOR (PG-LB)
                                                                                                                                                                                                          Signal; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Choi H:U.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 40; DB Pred. No. 27; 2; Mismatches
                       LRR
LRR
LRR
LRR
LRR
                                                                                                   LRR
                                                                                                               PROTEOGLYCAN.
7 LEUCINE-RICH TANDEM REPEATS
                                                                                                                                              SMALL CHONDROITIN/DERMATAN SULFATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C581EC33B8AC765A
                                                                                                                                                                                              matrix.
                                                                                                                                                                                                                                                                                                    TO
(INCOMPLETE).
                                                                                                   (INCOMPLETE).
                                                                                                                                                                                                                                                                                                    HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ruminantia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AND DISULFIDE BONDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Garza S., Hook
l proteoglycan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A
                                                                                                                                                                                                                                                                                                    OIF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14;
                                                                                                                                                                                                            Leucine-repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2;
                                                                                                                                                                                                                                                                                                                                  SEGMENT IS FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 1199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hook M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pecora;
                                                                                                                                                                                                                                                                                                                                                                                                                                       IN THE ZONE OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lugens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       THROUGH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ALSO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            reovirus:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a leucine-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Neame P.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MATRIX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q99645;
Q99645;
01-MAY-1997;
01-MAY-1997;
01-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DISULFID
DISULFID
DOMAIN
CARBOHYD
CARBOHYD
CARBOHYD
CARBOHYD
CARBOHYD
          DOMAIN
REPEAT
REPEAT
REPEAT
REPEAT
                                                                                           PROPEP
                                                                                                                                                                                               EMBL; U5911
MIM; 601657
                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A., AND TISSUE SPECIFICITY.
TISSUE-CHONDROCYTES;
MEDLINE-97131519; PubMed-8975717;
Deere M., Johnson J., Garza S., Harrison W.R., Yoon S.-J.,
Elder F.F.B., Kucherlapati R., Hook M., Hecht J.T.;
"Characterization of human DSPG3, a small dermatan sulfate
                                                                                                                             Connective
                                                                                                                                                   PFAM; PF00560; LRR; 3. PFAM; PF01462; LRRNT; 1.
                                                                                                                                                                                                                                                          ÷
                                                                                                                                                                                                                                                                                                                  Genomics 38:399-404(1996).

-I- FUNCTION: MAY HAVE A ROLE IN BONE FORMATION, AND ALSO IN
-I-STRABLISHING THE ORDERED STRUCTURE OF CARTILAGE THROUGH MATRIX

ORGANIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01 MAY-1997 (TYEMBLYEL 03, Created)
01-MAY-1997 (TYEMBLYEL 03, Last seq
01-CT-2000 (TYEMBLYEL 15, Last ann
SMALL CHONDROITIN/DERWATAN SULFATE P
                                                                                                                   SIGNAL
                                                                                                                                        Proteoglycan;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PGLB) (EPIPHYCAN)
DSPG3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9606
                                                                                                                                                                           INTERPRO;
                                                                                                                                                                                      NTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13 ILDAAVTAPTLE 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 VLNASVTSPLLE
                                                                                                                                                                                                                                  MANY PROTEINS. NUMBER IN THE SIMILARITY: TO OTHER SMALL
                                                                                                                                                                                                                                          SIMILARITY: THE REPEATED LEUCINE-RICH (LRR) SEGMENT IS FOUND MANY PROTEINS. NUMBER IN THIS PROTEIN: 6.
                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: EXTRACELLULAR MATRIX (BY SIMILARITY). TISSUE SPECIFICITY: CARTILAGE, LIGAMENT, AND PLACENTA. THE THREE GLYCOSAMINOGLYCAN CHAINS ARE DERMATAN SULFAI
                                                                                                                                                                                                                         PROTEOGLYCANS
                                                                                                                                                                                                                                                                    (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 7; Conser
                                                                                                                                                                          IPR000372; -. IPR001611; -.
                                                                                                                          tissue;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      117
278
77
. 60
. 64
95
282
282
319
321 AA;
153
168
192
212
238
259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                        Glycoprotein;
                                                                                                                                                                                                             AAC50945.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lrel. 03, Last sequence update)
Lrel. 15, Last annotation update)
DERMATAN SULFATE PROTEOGLYCAN PRECURSOR (PG-LB)
(DERMATAN SULFATE PROTEOGLYCAN 3) (DSPG3).
                                                                                                                             Extracel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    129
311
86
60
64
95
282
  301
167
191
191
211
237
238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        69.1%;
58.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       36687 MW;
                                                                                                                                      Signal; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROBABLE.
                                  LRR
LRR
LRR
                                                                     PROTEOGLYCAN
7 LEUCINE-PT
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GLCNAC
GLCNAC
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                                                                  LEUCINE-RICH TANDEM REPEATS
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                                                       (INCOMPLETE).
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(POTENTIAL).
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                                                                                                                                        Leucine-repeat;
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B S F F F

027116; 027116; 01-JAN-1998 01-JAN-1998 01-MAY-2000

(TrEMBLrel. (TrEMBLrel.

13,05,

Created)
Last sequence update)
Last annotation update)

PRELIMINARY;

PRT;

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Matches 7
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Best Local :
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CARBOHYD
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DISULFID
DISULFID
CARBOHYD
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CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., Oliver S., Osborne J., Quall M.A., Rajandream M.A., Rogers J., Rutter S. Seeger K. Skelton S. Schares S. Schares R. Steleton J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUN-1998 (TrEMBLrel.
01-JUN-1998 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
                                                                                                                                                                                                                                                                                 SEQUENCE
                                                                                                                                                                                                                                                                                                   INTERPRO; IPR000084;
PFAM; PF00934; PE; 1
PRODOM; PD001223; -;
                                                                                                                                                                                                                                                                                                                                                                                                          complete genome sequence. *;
Nature 393:537-544(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rutter S., Seeger K., Skelton S., Squares S., Sqares R., Sul
Taylor K., Whitehead S., Barrell B.G.;
"Deciphering the biology of Mycobacterium tuberculosis from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN-H37RV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria; Firmicutes; Actinobacteria; Actinobacteridae; Actinomycetales; Corynebacterineae; Mycobacteriaceae; Mycobacterium.NCBI_TaxID=1773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RV0742 OR MTV041.16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PGRS-FAMILY PROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=98295987; PubMed=9634230;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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                                                                                     96 ASITSPLLD
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                                                                                                           4 ASVTSPLLE
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77.8%;
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                                                                                                                                                                     Score 37; DB Pred. No. 15; 2; Mismatches
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Pred. No.
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GLCNAC (POTENTIAL).
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BY SIMILARITY.
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RESULT
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Q9LTR5
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J. Bacteriol. 179:7135-7155(1997).
EMBL; AE000876; AAB85533.1;
INTERPRO; IPRO00130;
IPROSITE: PS00142; ZINC_PROTEASE; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Harrison D., Hoang L., Keagle P., Lumm W., Pothier B., Qiu D., Spadafora R., Vicare R., Wang Y., Wierzbowski J., Gibson R., Jiwani N., Caruso A., Bush D., Safer H., Patwell D., Prabhakar McDougall S., Shimer G., Goyal A., Pietrovski S., Church G.M., Daniels C.J., Mao J.-I., Rice P., Nolling J., Reeve J.N.;
                                                                                                                                                                                                          Blaisonneau J., Nosek J., Fu
Submitted (MAR-1997) to the
EMBL; Y11606; CAA72340.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=DELTA H;
MEDLINE=98037514; PubMed=9371463;
                                                                                                                                                                                                                                               STRAIN=CBS 7907;
                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                            Saccharomycetaceae;
                                                                                                                                                                                                                                                                                                       Eukaryota; Fungi; Ascomycota;
                                                                                                                                                                                                                                                                                                                              Pichia kluyveri (Yeast).
                                                                                                                                                                                                                                                                                                                                            DNA POLYMERASE
                                                                                                                                                                                                                                                                                                                                                     01-JAN-1998 (TrEMBLrel. 01-MAY-2000 (TrEMBLrel.
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01-JAN-1998 (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                           Mitochondrion.
                                                                                                                                                                                  INTERPRO; IPRO02064; -.
PROSITE; PS00116; DNA_POLYMERASE_B; UNKNOWN_1
                                                                                                                                                                                                                                                                                 NCBI_TaxID=36015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical
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                                                                              VLNASV--TSPLLE 12
                                                         VLNASLRKTSPLFE
                                                                                                     ch 67.3%; l Similarity 71.4%; 10; Conservative
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PRELIMINARY;
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                                                                                                                                                              130725 MW;
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ne EMBL/GenBank/DDBJ
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Pred.
1; Mis
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Last
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Pred. No.
.PRT;
                                                                                                                                                                                                                                                                                                        Saccharomycetes;
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                                                                                                                                                               773C5756844E39A4 CRC64;
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1424 AA
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AC Q9
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Best Local S
Matches 7
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Best Local S
Matches 8
Q9NML8;
Q9NML8;
01-OCT-2000
01-OCT-2000
                                                                                                                                                                                   Murphy L., Quail M., Harris Submitted (JUL-2000) to the EMBL; AL160493; CAB97690.1; NON_TER 1 1
                                                                                                                                                                                                                                                                                                01-OCT-2000 (TREMBLIEL 15, 01-OCT-2000 (TREMBLIEL 15, 01-OCT-2000 (TREMBLIEL 15, PROBABLE ADA2-LIKE PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              features of the regions of (clones,");
DNA Res. 7:131-135(2000).
EMBL; AB024036; BAB02808.1;
SEQUENCE 1424 AA; 154737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
SIMILARITY TO BACTERICIDAL PERMEABILITY-INCREASING
                                                                                                                                                                                                                                                                   Leishmania major.
Eukaryota; Euglenozoa;
                                                                                                                                                                                                                                                                                                                                          Q9NMMO;
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Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9LTR5;
01-OCT-2000
                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                            STRAIN-FRIEDLIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=COLUMBIA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sato S., Nakamura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-COLUMBIA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arabidopsis thaliana (Mouse-ear cress).
                                                                                                                                                                                                                                                          NCBI_TaxID=5664;
                                                                                                                                                                                                                                                                                         LM26.72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Structural analysis of Arabidopsis thaliana chromosome 3. features of the regions of 4,504,864 bp covered by sixty P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nakamura
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                                                                                                                                                                                                                                       EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                       483 VLNMSVTSP 491
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                                                   11
                                                                                                                                                                                                                                                                                                                                                                                                                  1 VINASVTSP 9
                                                                                                    1 VLNASVTSPLLE
                                                                                  VLGANVTHPLMQ
                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Conserv
                                                                                                                       Similarity 58.37; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (FEB-1999)
                                                                                                                                                                           52
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                               PRELIMINARY;
                                                                                                                                                                           AA;
                                                                                  .40
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                                                                                                                                                                           5639 MW;
                                                                                                                                  65.5%;
58.3%;
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88.9%;
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to the E
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                                                                                                                                                                                                      D., Rajandream M., EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                            Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , WM
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                                                                                                                                                                                                                                                                                                                                Created)
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                                                                                                                        Score 36; DB Pred. No. 7; 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 37;
Pred. No.
                                                                                                                                                                          OBD2006CA2F21EC5 CRC64
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                                PRT;
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1.3e+02;
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                                                                                                                                            Length 52
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                                                                                                                         Indels
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e; eurosids II;
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(TrEMBLrel. 15, (TrEMBLrel. 15,

Created)
Last sequence update)

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RESULT
Q9UUST
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D7 0
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Best Local Similarity
---has 7; Conserv
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Best Local S
Matches 7
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
NCBI_TaxID=9606;
[1]
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01-AUG-1998 (Trem
01-JUN-2000 (Trem
IMPORTIN ALPHA 7
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Murphy L., Quail M., Harris
Submitted (JUL-2000) to the
EMBL; AL160493; CAB97692.1;
NON_TER 1 1
                                                                                                                  01-OCT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 15,
01-OCT-2000 (TrEMBLrel. 15,
01-OCT-2010 (TrEMBLREL. 15,
0162215:1 (IMPORTIN ALPHA 7
                                                                                                                                                                                Q9NU85;
01-OCT-2000
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                                                                                                                                                                                                                          Q9NU85
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PFAM; PF01749; SEQUENCE 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Christiansen M., Hart
Submitted (APR-1998)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leishmania major.
Eukaryota; Euglenozoa;
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                                                                              Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                            205 VLNCSILNPLL 215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRO; IPR002652; -. PF00514; Armadillo_seg;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AF060543; AAC15233.1; Q02821; 1BK5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 58.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                121; 1BK5.
IPR000225;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TremBLrel. 07, Created)
(TremBLrel. 07, Last sequence update)
(TremBLrel. 14, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
                                                                                                                                                                                                                          PRELIMINARY;
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AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hartmann E.;
998) to the EMBL/GenBank/DDBJ databases
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                                        Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chordata;
Primates;
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63.6%;
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58.3%;
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                                                                                                                    Created)
Last sequence update)
Last annotation update)
(KARYOPHERIN))
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                                                                                                                                                                                                                                                                                                                                                                                                                              Score 36;
Pred. No.
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Pred. No.
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                                      Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata;
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                                                                                                                                                                                                                          566
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74;
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18;
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Best Local S
Matches 7
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Best Local S
Matches 7
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01-MAY-1997
01-MAY-2000
                                                                                                                        P87883;
01-MAY-1997
01-MAY-1997
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (FEB-2000) to the EMBL/GenBank/DDBJ EMBL; AL049795; CAB75614.1; - SEQUENCE 566 AA; 63004 MW; 48657248DFF2678
                                                                                                                                                                                                                                                                                                                                                               epitope of hepatitis C virus in two haemophilic patients. J. Gen. Virol. 78:583-590(1997). EMBL; Z84308; CAB06380.1; -.~
         STRAIN-PATIENT 1, 1994;
MEDLINE-97201623; PubMed-9049409;
Devereux H.L., Brown D., Dusheiko
                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-PATIENT 2, 1989;
MEDLINE-97201623; PubMed-9049409;
Devereux H.L., Brown D., Dusheiko
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Viruses;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                         SEQUENCE FROM N.A.
                                                                      Viruses; ssRNA positive-strand Hepacivirus
                                                                                                      NS4.
                                                                                                               PARTIAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P87859
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                                                                                                                                                                  P87883
                                                                                                                                                                                                                                                                                                               SEQUENCE
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                                                            NCBI_TaxID=11103;
                                                                                           Hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=11103;
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 .L., Brown evolution
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7 (TrEMBLTel. 03, I
0 (TrEMBLTel. 13, I
1 GENE (PATIENT 1,
                                                                                                                                                                                                                                                                                                               44 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (Trembirel. 03, (Trembirel. 03, I) (Trembirel. 13, I) (Trembirel. 13, I) GENE (PATIENT 2,
                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                  PRELIMINARY;
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 D., Dusheiko of the 5'UTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.5%;
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he 5'UTR and a region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Created)
Last sequence update)
Last annotation update)
, 1989, CLONE 4) (FRAGMENT).
                                                                                                             Last sequence update)
Last annotation updat
, 1994, CLONE 3) (FRAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              <u>ب</u>
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Pred.
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                                                                                                                                                                   PRT;
                                                                                 viruses,
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G.M.,
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79;
                                                                                                                                                                                                                                                                    ).3;
                                                                                 no
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Emery V.C., Le
region of NS4
                                                                                                                                                                                                                                                                                                                                                                                                          Emery V.C.,
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                                                                                  DNA
                                                                                                                                                                                                                                                                                 14;
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                                                                                                              update)
(FRAGMENT)
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                                                                                 stage;
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           Lee
                                                                                                                                                                                                                                                                                                                                                                                                          Lee C.A.;
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  containing
                                                                                   Flaviviridae;
           C
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Search completed: June 4, 2001, 12:23:21 Job time: 584 sec À Query Match 63.6%; Score 35; DB 14; Length 44; Best Local Similarity 77.8%; Pred. No. 9.3; Matches 7; Conservative 1; Mismatches 1; Indels epitope of hepatitis C virus in two haemophilic patients.";
J. Gen. Virol. 78:583-590(1997).
EMBL; Z84361; CAB06413.1; -INTERPRO; IPR000745; -PFAM; PF01006; HCV\_NS4a; 1.
NON\_TER
NON\_TER
1 1 1
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44 44
5EQUENCE 44 AA; 4317 MW; 71D5F7858B3A670E CRC64; 2 LNASVTSPL 10 | ||:|||| |4 LTASITSPL 12 0;

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Result
No.
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Maximum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                          Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                seq
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length: 2000000000
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61
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Gapop 10.0 , Gapext.0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /SIDS6/gcgdata/geneseq/geneseqp/AA1980.DAT:*
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/SIDS6/gcgdata/geneseq/geneseqp/AA1999.DAT:
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Y33183
                                                                                                  196183
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q/geneseqp/AA1994.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14 . DAT :
                                                                                                                                                                                          Description
                                                                                                                                                           Oligopeptide from
                                                 Human
                                                                 Human
                                                                                Human macrophage-s
                                                                                             Human macrophage-s
                                                                                                               Human
                                                                                                                            Human
                               n macrophage-s
n FcRI (cDNA c
n FcRI (cDNA c
                                                                                                               FCRI (CDNA C
                                                                                                                            macrophage-s
                                                                                                                                              sfc-gammaRI
receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
W60554
ID W6
XX
AC W6
XX
Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                             W60554;
                                                                             WPI; 1998-240964/21.
                                                                                                                                                                          08-OCT-1996;
                                                                                                                                                                                                         07-OCT-1997;
                                                                                                                                                                                                                                                                      WO9815833-A1
                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                              18-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W60554 standard; peptide; 12 AA
                                                                                                          De Kruif CA, Logtenberg
                                                                                                                                           (UYUT-) RIJKSUNIV UTRECHT.
                                                                                                                                                                                                                                        16-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                          96EP-0202791
                                                                                                                                                                                                         97WO-NL00557.
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## ALIGNMENTS

Oligopeptide from extracellular domain of CD64 immunohistochemical; immunofluorescer
cell transformation; mutation; anti; Extracellular domain; CD64; identification; antibody; immunofluorescent analysis; detection;

surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms. Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated affinity binding to the CD64-covered rods. The specification describe forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode method for the identification of a peptide able to bind specifically to target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused Example 1; Page 29; 40pp; English. Sequence by mutation in 12 ΑĄ, (anti)oncogenes The specification describes isolated by þ

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                                                                                                                                                          Query Match
Best Local S
Matches 12
Human sFc-gammaRI protein fragment.
                                                               Y33183 standard;
                      15-NOV-1999
                                                                                   N
                                                                                                                gnlvtlscetkl 12
                                                                                                                                       GNLVTLSCETKL 12
                                                                                                                                                           l Similarity
12; Conser
                                                                                                                                                          100.0%; llarity 100.0%; Conservative
                     (first entry)
                                                               Protein;
                                                               261
                                                                                                                                                            0;
                                                                                                                                                                     Score 61; DB 19;
Pred. No. 0.00032;
                                                               AA.
                                                                                                                                                            Mismatches
                                                                                                                                                       Indels
                                                                                                                                                                               Length 12;
                                                                                                                                                             0;
                                                                                                                                                            Gaps
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anti-inflammatory; drug design; therapy; inhibitor; tissue damage; atomic Fc receptor; Fc-gammaRIIa; human; FCR; nflammatory coordinate; modulator; Fc-gammaRIa; hypersensitivity; bioactive compound design; computer-assisted; py; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; Ig 'rc-gammaR; Fc-epsilonR; IgG; IgE; inflammatory cell recruitment; immune function regulation; model; three-dimension; 3-D;

## Homo sapiens

12-AUG-1999

04-FEB-1999; 99WO-IB00367

11-SEP-1998; 06-FEB-1998; 98US-0099994. 98US-0073972.

(ILEX-) ILEXUS PTY LTD

Baell JB, Maxwell KF, Epa V, Garrett TPJ, Hogarth PM, McCarthy TD, McKenzie IFC, Pi Pietersz GA; Matthews BR

WPI; 1999-539978/45

Three-dimensional structures and models of Fc receptors, useful computer-assisted drug design  $% \left\{ 1\right\} =\left\{ 1$ 

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Claim 11; Page 316-317; 326pp; English
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hypersensitivity recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc gamma RIa and models of other FCR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein oligonucleotides, the FcR can be used to design bioactive chemical fragment described reduce IgG- or IgE-mediated tissue damage, e.g. that e FCR can be used to design bioactive chemical compounds, e.g. igonucleotides, peptides, peptidomimetics and small organic molecules, computer-assisted methods of drug design. Therapeutic compositions at inhibit the activity of FC-gammaR or FC-epsilonR can be used to in the method of the IgG- or IgE-mediated ç

Sequence

Query Match Best Local 9 Matches Local Similarity nes 12; Conserv Conservative 100.0%; 0; Score 61; DB 20; Pred. No. 0.0073; Length 261; Indels 0; Gaps

0;

1 GNLVTLSCETKL 12

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Qy

RESULT R22549

R22549 standard; Protein; 344

21-MAY-1992 (first entry)

Human macrophage-specific FcRI receptor encoded by clone p98

Rapid immunoselection cloning technique; immunodiagnosis; high affinity receptor. technique; cell surface antigen;

Homo sapiens

W09201049-A

XX PX XX

PD XX PR XX PR XX PR XX PR XX PR XX PR

15-JUL-1990; 90WO-US04986

13-JUL-1990; 90US-0553759

(GEHO-) GEN HOSPITAL CORP

Seed B, Aruffo A, Amiot M;

N-PSDB; WPI; 1992-056864/07 N-PSDB; Q21179.

New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic it - for
neoplasms,

Example 10; Page 94a; 160pp; English

CCCXXXTTXR This amino acid sequence was predicted from the cDNA sequence of cDNA clone 998/X2. It differs from the sequence predicted from clone 9135 (see 021178) at position 25; 9135 encodes a Ser residu while clone 998/X2 predicts a Thr residue. At position 58, 990Ser residue 58, p90

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Query Match
Best Local S
Matches 12
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13-JUL-1990;
18-OCT-1993;
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macrophages. 2 Other cDNA ciones, pio (1 coded for a variants of the FCRI sequence; of the p98 product is truncated compared w
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
           for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FcRI sequence; the C-terminal sequence of the p98 product is truncated compared with those of the p135
                                                                                                    cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity reco
                                                                                                                                                                                                                                                           Cloning of cDNA encoding cell surface antigen -
                                                                                                                                                                                                                                                                                                        N-PSDB; T14718
                                                                                                                                                                                                                                                                                                                                                                           Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cell surface antigen; therapy; diagnosis; ve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R91439 standard; Protein; 344 AA.
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                                                                                                                                           The amino acid sequence (R91439) of human FcRI was detd. from a
                                                                                                                                                                                        Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               205
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90US-0553759.
93US-0139273.
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88US-0160416
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ector; FcRI; Fc receptor.
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translated
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1 product"
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                                                                                                                                                                                                                                                              useful
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                                                                                                    receptor
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Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and p90 products (see also R91438 and W00859). A of FCRI and a receptor ligand will be helpful in potency of antibodies in therapy.
           Misc-difference
                                                Misc-difference
                                                                    Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                            panning; immunodiagnosis; diagnosis; immunotherapy;
immune disorder; infection; asthma; immune-complex of
amyloidosis; multiple sclerosis; parasitic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage-specific FcRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Y96183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y96183 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Macrophage; FcRI; cell surface antigen; human;
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9 51
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136
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117
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122
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139 ·
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                                                                                                                                                                                                                                                                                                                                                                                                                                  immunotherapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                            immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A fusion protein
In increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 344;
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В

205 gnlvtlscetkl 216

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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiting accomplex disease, parasiting accomplex disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-586382/55.
N-PSDB; A50631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
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surfaces other than macrophages, protein of FCRI and a receptor 1
                                       sclerosis. FCRI is a high affinity receptor for the FC poligg, normally located on the cell surfaces of macrophages ability to interfere with such bonding, or to cause it to
                                                                                                                                                                                                                      physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                       physical selection of cells expressing the antigen by adhesion to
                                                                                                                                                                                                                                                                                                                                                           The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50631), which was isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13-JUL-1990
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                                                                                                                                                                                                                                                                                                                                          from
                                                                                                                                                                                                                                                                                                                                                                                                                              Example 10; Column 53-55;
                                                                                                                                                       immunotherapeutic applications, including treatment of immune-mediated infections, c
                                                                                                                                                                                                      proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                   the invention designed to isolate CSA nucleic acids.
                                                                   immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The
                                                                                                                                                                                                                                                                                                                                          uced from cDNA clone p135 (see A50631), which was isolated cDNA library expressed in COS cells using a novel method of
                                                                                                                                including humans.
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306
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88US-0160416
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216
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333
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305
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268
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220
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338
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332
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                                                                                                                                                                                                                                                                                                                                                                                                                            75pp;
                                                                                                                                    These disorders include asthma,
bonding, or to cause it to occur on , is useful in therapy. A fusion ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         bу
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                                                                                                                                                       diseases,
                                                                                                                                                                              the diagnosis and
                                                                                                                                                         and disorders in
                                                                                                                                                                                                                                                                                                                        The method is
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Matches

1 GNLVTLSCETKL 12

Query Match
Best Local Similarity

100.0%;

Score 61; DB 21; Pred. No. 0.0097;

DB 21;

Length 344; Indels

0;

Mismatches

0;

0

Gaps

21-MAY-1992

(first entry)

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Sequence

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                                                                                                                                                                                                                                         Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25, p135 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the predicts Val. Sequences predicted from all 3 clones show the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R20811 standard; Protein;
                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                 potency of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10; Page 94a; 160pp; English:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO-) GEN HOSPITAL CORP
                                                                                   R22550 standard; Protein; 374 AA
                                                                                                                                                                       205
                                                                                                                                                                                                        1 GNLVTLSCETKL 12
                                                                                                                                                                       gnlvtlscetkl
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                                                                                                                                                                                                                                                                                                                                    374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90WO-US04986
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                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                               Score 61; DB 13
Pred. No. 0.011;
                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                 DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ıt.
                                                                                                                                                                                                                                                                               Length 374;
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Best Local Similarity
Matches 12; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane spanning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residu
                                                                                                                  Cell surface antigen; cloning; immunoselection; immunotherapy;
therapy; diagnosis; vector; FcRI; Fc receptor.'
                                                                                                                                                                                            30-OCT-1996
                                                                                                                                                                                                                                                    R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and clone p90 predicts a Thr residue. At position 50, p90 predicts Val while clones p135 and p98/X2 (see Q21179) both predict Leu.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
Misc-difference
                                                                                        Homo sapiens
                                                                                                                                                               Human FcRI (cDNA clone pl35 product).
                                                                                                                                                                                                                         R91438
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-JAN-1992
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                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; Page 94a; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human macrophage-specific FcRI receptor encoded by clone p90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                           205 gnlvtlscetkl 216
                                                                                                                                                                                                                                                                                                                                                        1 GNLVTLSCETKL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             374 AA;
                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90US-0553759
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                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   It differs from the sequence predicted from 021178) at position 25; pl35 encodes a Ser residue
                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 61; DB 13
100.0%; Pred. No. 0.011;
tive 0; Mismatches
               "amino acid 25 is Thr in
translated products"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English
                                                                                                                                                                                                                                                                                                                                                                                                               DB 13;
                                                                                                                                                                                                                                                                                                                                                                                     0
                             clone p90 and p98
                                                                                                                                                                                                                                                                                                                                                                                                             Length 374;
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Best Local Similarity
Matches 12; Conserv
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                       09-APR-1996
                                                                                                                    US5506126-A
                                                                                                                                                                           Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FCRI; FC receptor.
                                                                                                                                                                                                                      Human FcRI (cDNA clone p90 product)
                                                                                                                                                                                                                                                    30-OCT-1996
                                                                                                                                                                                                                                                                                                           W00859 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10; Column 55-56; 79pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-APR-1996
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                                                          25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHO ) GEN HOSPITAL CORP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
                92US-0983647.
88US-0160416.
                                                           88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88US-0160416
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 61;
Pred. No.
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cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (#00859) of the FCRI sequence, and a third clone, p98 (T14718), coded for an FCRI (#91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cloning of cDNA encoding cell surface antigen - useful of diagnostic and therapeutic proteins % \left( 1\right) =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1\right\} =\left\{ 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "amino acid 58 is Val in p90 clone translated product"
                     Mismatches
                                                                                                                                                                                              0.011;
                                                                                                                                                                                                                                                                                                                                                                                                                  DB 17;
                     0
                                                                                                                                                                                                                                                                                                                                                                                             Length 374;
                     Indels
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                               /note=
Misc-difference 85
                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The amino acid sequence (W00859), of human FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity recept for the FC portion of IgG, normally located on cell surfaces of macrophages. Other CDNA clones (see also T14717-18) coded for macrophages.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
         Misc-difference
                               Misc-difference
                                                         Misc-difference
                                                                                                                                  Misc-difference
                                                                                                                                                           Misc-difference
                                                                                                                                                                              /note=
Misc-difference 45
                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                             Human Fc receptor I
                                                                                                                                                                                                                                                                                                                                                                                                             W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                   W80448 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    variants (R91438-39) of the FCRI sequence. FCRI and a receptor ligand will be helpful potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                          cloning
                                                                                                                                                                                                                                                                                                                                    Fc receptor I; FcRI; cell surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; T14719
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                              Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GNLVTLSCETKL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                     /note=
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93US-0139273
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159
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99
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23
                                                                                                                                                                                                                                                                Location/Qualifiers
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                                            'note=
                                                                     'note=
                                                                                                                                                                                                                     'note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                              "encoded
                                                                   "encoded
                                                                                                                                            "encoded by CCC"
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                encoded by
                                         "encoded by CCC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 61; DB 17; Length 374; Pred. No. 0.011;
                                                                  by
                                                                                                                    уd
                                                                                                                                                                     by CTG"
                                                                                           by TCC"
                                                                                                                                                                                               by GAC"
                                                                                                                                                                                                                       by CTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                   antigen; lymphocyte; human;
                    GGC"
                                                                     CAA"
                                                                                                                      AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A fusion protein of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT W97833

11

AC XX

W97833 standard; Protein; 374 AA.

W97833;

g

205 gnlvtlscetkl 216

1 GNLVTLSCETKL 12

Qy

Query Match 100.0%; Best Local Similarity 100.0%; Matches 12; Conservative

0

Score 61; DB 19 Pred. No. 0.011; Mismatches

19; 0;

Length 374; Indels

0

Gaps

0,

Sequence

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13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                               transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V6342-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note=
Misc-difference 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
Misc-difference_176
                                                                                                                                                                                                                                                                                                                                                                       in eukaryotic cells and selection of cells expressing the antigen be adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone pl35 (see V63456) isolated using a rap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see x07372) and p98 (see x07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human lymphocyte antigenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-1992;
25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoselection cloning method from a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1998-609251/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The cDNA library was constructed
                                                         including humans.
374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                 obtained. A novel method for cloning cDNAs from mammalian libraries is based on transient expression of an antigen by tic cells and selection of cells expressing the antigen by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89US-0379076
90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0861205
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88US-0160416
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171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "encoded by AAC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bу
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTC"
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This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in C cells. The cDNA library was constructed from polya RNA of cells from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mammal
                                                                                                                                                             New cloning vector and polylinker - based on existing sequences fefficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                            13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
Misc-difference 256
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Misc-difference 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "encoded Misc-difference 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "encoded by TCC" Misc-difference 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=
Misc-difference 85
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N-PSDB; X07372.
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25-FEB-1988;
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                                                                                                                               Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                                            Aruffo A,
                                                                                                                                                                                                                                                                                                                                             21-MAY-1997
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? 77
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89US-0379076.
90US-0553759.
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103
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176
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expression libraries

is based on transient expression of an antigen

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W97834;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W97834 standard; Protein;
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                                   Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                Human Fc receptor I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    animals, including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunotherapeutic applications, includin treatment of immune-mediated infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the generation of very large mammalian expression libraries. purified genes and proteins are useful for immunodiagnostic a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        genes can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the isolation and cloning of any protein which can be expressed and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 205 gnlvtlscetkl 216
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The method
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RESULT 13
Y96134
ID Y96134
XX
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AC Y96134
XY
DT 19-DEC
XX
DT 19-DEC
XX
Human
XX
Macrop
KW Macrop
KW Manune
KW immune
KW amyloi
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                                                                                                                                                                                                                                                                                                                                                                                                                                expression libraries is based on transient expression of an antigen to end to end to end to end to end to end to end to end to end to end to expression of an antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see W63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunother apeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 12; Conserv
                         Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therap immune disorder; infection; asthma; immune-complex disease;
            panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
                                                                                    Human macrophage-specific FcRI
                                                                                                                                                                         Y96134 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                   19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                           animals,
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25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Column 53-54; 75pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                          treatment of immune-mediated infections,
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                                                                                                                                                                                                                                                                                                                                                                               374 AA;
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89US-0379076.
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                                                                                                                                                                           374 AA
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Pred. No. 0.011;
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                                                                                                                                                                                                                                                                                                          0
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13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
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Misc-difference 129
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Misc-difference 126 ;
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Misc-difference 116
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89US-0379076.
90US-0498809.
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216
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332
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2000-586382/55

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Y96226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                      Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IGG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodlagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
                                                                           03-DEC-1998;
                                                                                                                                                      07-JUN-2000
                                                                                                                                                                                            EP1006183-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y96226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 53-55; 75pp; English
WPI; 2000-367968/32
                                                                                                               03-DEC-1998;
                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                     systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                       Human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Y96226 standard; Protein; 374 AA.
                                    (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          205
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lmmune-complex disease, amyloidosis, parasitic diseases or multiple
sclerosis. FcRI is a high affinity receptor for the Fc portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of immune-mediated infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mmunotherapeutic applications, including the diagnosis and isorders in reatment of immune-mediated infections, diseases, and disorders in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GNLVTLSCETKL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gnlvtlscetkl 216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                       erythematosus; tumour
                                                                             98EP-0122969.
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Pred. No. 0.011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of pattents with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds cells expressing Fc receptors. FcgammaRI molecules are expressed by
                                                prevent autoimmune diseases, allergies or tumours, especially AIDS rheumatoid arthritis or MM.
                                                                                            myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or
                                                                                                                                                                                                                                                                    may be modified to produce recombinant versions.
                                                                                                                                                                                                                                                                                           neutrophils and eosinophils. Upon Fc receptor activation, immune effecton pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                         monocytes and macrophages, but expression can also be induced
                                                                                                                                                                                                                                                                                                                                                                                                                                           regammaRI is also known as CD64. Fe receptors play an importan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel Fc receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the human high affinity Fc receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English
                                                                                                                                                                                                                                                                         The recombinant Fc
                                                                                                                                                                                                                                                                                                                          immune effector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FcgammaRI
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Query Match Best Local Matches Local 1 GNLVTLSCETKL 12 Similarity 100. 12; Conservative 100.0%; 0 Score 61; Pred. No. Mismatches 0.011; DB 21; 0; Length 374; Indels 0 Gaps

0

Sequence

374

AA;

밁 δÃ

B43683 ID B XX RESULT Human cancer associated protein sequence SEQ ID NO:1128. 08-FEB-2001 B43683 standard; Protein; 399 AA. 15 (first entry)

vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation; immune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection; neurological disease; drug screening. antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; diagnosis; dermatological; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; Human; cancer associated gene; cytostatic; neuroprotective; proliferative; vulnerary; cancer antigen; detection; thrombolytic; coagulant; nootropic; ımmunomodulator; cancer;

Homo sapiens

XSX WO200055350-A1

X P X P X P 21-SEP-2000

08-MAR-2000; 2000WO-US05882

12-MAR-1999; 99US-0124270.

(HUMA-) HUMAN GENOME SCI INC

Rosen CA; Ruben SM;

WPI; 2000-587533/55

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N-PSDB; C7789
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Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer  $\boldsymbol{\cdot}$ 

Claim 11; Page 1739-1740; 2352pp; English.

C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antitasthmatic; antirheumatic; antiarthritic; antitiric; antidiabetic; antitallergic; antibacterial; antiviral; dermatological; neuroprotective; cardiant; thrombolytic; coagulant; notropic; antipsoriatic and antianglogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and B44240 represent sequences used in the exemplification of the present invention.

Sequence 399 AA;

Query Match 100.0%; Score 61; DB 21; Length 399; Best Local Similarity 100.0%; Pred. No. 0.011; Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps

0;

Search completed: June 4, 2001, 12:13:27 Job time: 200 sec

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Perfect score:
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                            Score
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Match
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1: /cgn2_6/ptodata/:
2: /cgn2_6/ptodata/:
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Gapop 10.0 , Gapext 0.5
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61
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                                                                                              GenCore version
Copyright (c) 1993 - 2000
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US-08-530-146-16
US-08-985-950-14
US-08-985-950-14
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; ;		•			Sequence 25, Sequence 5, Sequence 5, Sequence 1, Sequence 2, Sequence 1, Sequence 1,

78.78;

Score 48;

DB 2;

Length 91;

75.0%;

Pred. No. 0.13;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens -08-297-395-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PPLICANT: Howard L. Weiner PPLICANT: David A. Hafler
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ARLIER APPLICATION NUMBER: 07/065,734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ARLIER APPLICATION NUMBER: 08/059,189
ARLIER FILING DATE: 1993-05-06
ARLIER APPLICATION NUMBER: 07/502,559
ARLIER FILING DATE: 1990-03-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ARLIER APPLICATION NUMBER: PCT/US88/02139
ARLIER FILING DATE: 1988-06-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  JRRENT FILING DATE: 1994-08-1
                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,860
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                                                                                                                                                                                                                                                                                                                                             PPLICANT:
                                                                                                                                                                                                                                               ORRESPONDENCE ADDRESS:
                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                   ITLE OF INVENTION: VACCINATION AND METHODS AGAINST DISEASES
ITLE OF INVENTION: RESULTING FROM PATHOGENIC RESPONSES BY SPECIFIC T CELL
ITLE OF INVENTION: POPULATIONS
                                                                                                                                                                       CITY: SAN DIEGO
STATE: CALIFORN
                                                                                                                                                                                                                                                                     JMBER OF SEQUENCES:
                                                                                                                                      COUNTRY: 1
                                                                                                                                                                                                            STREET:
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|7 GSLVTLNCETNL 28
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                                                                                                                                                  CALIFORNIA: UNITED STATES
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                                                                                                                                                                                                        E: CAMPBELL AND FLORES 4370 LA JOLLA VILLAGE.
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Best Local
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                 TELEFAX: 619-535-8949
NFORMATION FOR SEQ ID NO:
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ATTORNEY/AGENT INFORMATION
NAME: CAMPBELL, CATHRYN
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TELEPHONE: 619-535-9001
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PRIOR APPLICATION DATA:
                                                                            SEQUENCE CHARACTERISTICS: LENGTH: 113 amino acid
                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 619-535-9001
                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION: NAME: CAMPBELL, CATHRYN
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                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING TOTAL PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: SAN DIEGO
STATE: CALIFORNIA
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REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                             SOFTWARE:
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                                          TOPOLOGY:
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                                                            amino acid
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4370 LA JOLLA VILLAGE DRIVE, SUITE 700
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                                                                               113 amino acids
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                                          linear
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N-terminal
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RESULTING FROM PATHOGENIC RESPONSES BY SPECIFIC T CELL
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Pred. No. 13;
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Length 113; Indels `

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Length 113;

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Best Local Similarity bu...
Thes 8; Conservative
             Sequence 14, Application US/08442542 Patent No. 5686600
                                                                                                                                                            Query Match
Best Local Similarity
Matches 8; Conser
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GENERAL INFORMATION:
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NFORMATION FOR SEQ ID NO:
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LENGTH: 113 amino acid
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PRIOR APPLICATION DATA:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                           amino acid
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CARLO, DENNIS J.
CARLO, DENNIS J.
VENTION: VACCINATION AND METHODS AGAINST DISEASES
VENTION: RESULTING FROM PATHOGENIC RESPONSES BY SPECIFIC T CELL
                                                                                                                                                            Conservative
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N-terminal
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Pred. No.
                                                                                                                                                                          Score 37;
Pred. No.
                                                                                                                                                                                       DB 4;
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Query Match
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                                                                                                                                                                                                                                                                                                         tent No.
                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
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            SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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MEDIUM TYPE: Floppy disk
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                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Palana
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                                                                                                                     COUNTRY:
                                                                                                                                                  CITY: Hawthorne
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                                                                                                                       USA
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SYSTEM: PC-DOS/MS-DOS
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70.0%;
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US/08/765,469
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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Best Local Similarity
Thes 7; Conserv
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US-08-765-469-14
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -US95-15696-2
                                                         TELEFAX: 207-363-0528
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 919-541-8689
NFORMATION FOR SEQ ID NO: '14:
SEQUENCE CHARACTERISTICS:
                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 207-363-0558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: President and Fellows of Harvard College TITLE OF INVENTION: SINGLE CHAIN T-CELL RECEPTOR NUMBER OF SEQUENCES: 14
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Spruill, W. Murray
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                 TTTORNEY/AGENT INFORMATION:
NAME: Farrell, Kevin M;
REGISTRATION NUMBER: 35,505
REFERENCE/DOCKET NUMBER: HU
                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: CGITELECOMMUNICATION INFORMATION: TELEPHONE: 919-541-8615
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                         TYPE: amino acid
                 TOPOLOGY:
                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/US95/15696
                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: York Harbor
                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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                                             391 amino acids
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                                                                                                                                                                                                                                                                                                                                                              Floppy disk
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Pred. No. 14;
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                                                   Matches
                                                          Query Match
Best Local Similarity
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Patent No.
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Best Local
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                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                          IMMEDIATE SOURCE:
LIBRARY: GenBa
CLONE: 969078
                                                                                                                                                                                                                                                          TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/8 FILING DATE: Filed Herewith PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
403 GRLVNLSCST 412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ITLE OF INVENTION:
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                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                          TYPE: amino acid
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                        1 GNLVTLSCET 10
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5. 5854023
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8; Conserv
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                                                  Conservative
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                                                                                                                                         GenBank
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                                                                                                                                                                  linear
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                                                             59.0%;
70.0%;
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                                                             Score 36; DE
Pred. No. 91;
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                                                                          Length 504;
                                               Indels
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RESULT 10 US-08-037-816A-20

Sequence 20,

Application US/08037816A

Patent No. 5869624
GENERAL INFORMATION:
APPLICANT: Hasel,
APPLICANT: Maddon,

Hasel, Karl W. Maddon, Paul J.

TITLE OF INVENTION: PITLE OF INVENTION:

THEREOF

HIV-1 VACCINES, ANTIBODY COMPOSITIONS RELATED THERETO, AND THERAPBUTIC AND PROPHYLACTIC USES

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Query Match
Best Local Similarity
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TELEX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 20
SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acids
TYPE: amino acid
         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.24
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/037,816A
FILING DATE: 26-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                    equence 24, Appliatent No. 5869624
                                                                                                                                                                                                                                                                                                                                                                                                                                     08-037-816A-24
                                                                                                                                                                                                                                                                                                                                                                                  ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERRICE/DOCKET NUMBER: 41190/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEPHONE: (212) 977-9550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Relaction CATA:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
Cooper & Dunham
                                                                                                                                                                                                                                                                                                FITLE OF INVENTION:
FITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                    CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 20
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/037,816A FILING DATE: 26-MAR-1993
CLASSIFICATION:
                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                          STREET: '30 Rockefeller Plaza
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                                                                                                                                                                                                                                                                                            HIV-1 VACCINES, ANTIBODY COMPOSITIONS RELATED THERETO, AND THERAPEUTIC AND PROPHYLACTIC USES THEREOF
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Query Match
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"Thes 6; Conserv?
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                                 TELEFAX: (212) 977-9550
TELEX: 42523 COOPUI
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEFAX: (212) 664-0525
TELEFX: 422523 COOPUI
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                atent No.
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                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 411
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PM PC COMPANS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.24
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/037,816
FILING DATE: 26-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: White, John P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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CITY: New York
STATE: New York
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                TOPOLOGY:
                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 10112
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GY: linear
                               amino acid
TYPE:
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Maddon, Paul J.
VENTION: HIV-1 VACCINES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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54.5%;
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Pred. No. 1.3e+02;
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US-08-530-146-24
                                                                        Sequence 16, Application US/08037816A Patent No. 5869624
                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein -08-530-146-24
                                                          GENERAL INFORMATION:
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equence 24, Approper No. 5886163
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TELEX: 422523 COOPUI
NFORMATION FOR SEQ ID NO: 2
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1 24
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INVENTION: HIV-1 VACCINES, ANTIBODY COMPOSITIONS RELATED
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US-08-037-816A-16
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Best Local Similarity 54.
Matches 6; Conservative
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                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/037,816A
FILING DATE: 26-MAR-1993
                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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LENGTH: 507 amino acids
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TTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                            ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      408 GNTITLPCRIK 418
                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.24
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STATE: New York
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REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 41190/JPW/AJM
REFERENCE/DOCKET NUMBER: 41190/JPW/AJM
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEPHONE: (212) 644-0525
TELEX: 42253 CCOPUI
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 507 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: protein
US-08-037-816A-28

Ouery Match
Best Local Similarity 54.5%; Pred. No: 1.4e+02;
Matches 6; Conservative 1; Mismatches 4; Indels 0; Gaps
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## ALIGNMENTS

R;Allen, J.M.; Seed, B. Science 243, 378-381, 1989
Science 243, 378-381, 1989
A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Title: number: A41357; MUID:89100284
A;Accession: A41357

A; Molecule type: mRNA

C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000
C;Accession: A41357; S03019

Fc gamma (IgG) rece N;Alternate names:

receptor I (high affinity) form b

A41357 RESULT

A; Reference number: A39878; A; Accession: A39878

A; Residues: 1-374 < VAN> A; Molecule type: DNA

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A; Title: Linkage on chromosome 3 of autoin A; Reference number: 148471; MUID: 93242399 A; Accession: 148471
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Science 260,
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:Perez, C.; Wietzerbin, J.; Benech, P.D.
al. Cell. Biol. 13, 2182-2192, 1993
                                                          Status: preliminary; translated from
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Status: translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Superfamily: Fc gamma receptor I; immunoglobulin homology Keywords: alternative splicing; glycoprotein; immunoglobu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cross-references: GB:S57204; NID:g298692;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE type: DNA
Residues: 1-8 <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             293-313/Domain: transmēmbrane #status predicted <TMM>59,78,152,159,163,195,240/Binding site: carbohydrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: GDB:135911; OMIM:146760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOIECULE type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fitle: Nucleotide sequence Reference number: S03018; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Title: Isolation and expression of functional high-affinity Fc receptor complementary Reference number: A41357; MUID:89100284
Accession: B41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cross-references: GB:L03418;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Residues: 1-374 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          leic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ccession: 170304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6-292/Domain: extracellular #status predicted <EXT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ence 243,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-170/Domain: immunoglobulin homology <IMM2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             esidues: 1-24,'S',26-337,'T',339-374 <AL1>
coss-references: GB:X14356; GB:M21091; NID:g31331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cession: S03018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         lecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5/Domain: signal sequence #status
                                                                                                                                                                                                                                                                                                                                               205 GNLVTLSCETKL 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                     1 GNLVTLSCETKL 12
                                                                                                                                                                 J.В.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            А.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
12; Conser
                                                                                                                                          695-698, 1993
                                                                                                                                                                                    I48471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         378-381,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cis-DNA elements involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seed,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fc gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
ilarity 100.0%;
Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1q21-1q21
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90, 2102-2109, 1992
<RES>
s: EMBL:X70980;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  I55577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GB/EMBL/DDBJ
                                                                                                                                                           Rodrigues, N.R.; Ghosh, S.; Hogarth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e of three cDNAs
MUID:89098339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               eptor I family gene products MUID:93055454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GB:M63835; NID:g180277; PIDN:AAA35678.1; PID:g180279 Doebele, R.; Pan, L.C.; Salmon, J.E.; Kimberly, R.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NID: 9184840;
NID: 9311748; PIDN: CAA50311.1;
                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 61; DB
Pred. No. 0.0
); Mismatches
                                                                                                                         autoimmune diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             site: carbohydrate (Asn)
                                                              GB/EMBL/DDBJ
                                                                                                                                                                                                   02-Jul-1996 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            myeloid-cell-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       for
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                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
0.0022;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIDN: AAD13887.1; PID: g4261587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the human high affinity Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIDN:CAA32537.1;
                                                                                                                     and defective Fc
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PID:g311749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptor; transmembrane
                                                                                                                                                               P.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression
                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                               Wicker,
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                                                                                                                     receptor
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                                                                                                                                                                                                   C;Accession: A71890
R;Alm, R.A.; Ling, L.S.L.; Moir, Jives, C.; Gibson' R.; Merberg, Nature 397, 176-180, 1999
                                                                                                                                                                                                                                                                                hydroxyethylthiazole kinase - Helicobacter pylori (strain J99)
C;Species: Helicobacter pylori
A;Variety: strain J99
C;Date: 12-Feb-1999 #sequence_revision 12-Feb-1999 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
                                         A; Experimental source:
                                                                               A; Residues:
                                                                                                                                                             A; Title: Genomic sequence comparison of two unrelated isolates A; Reference number: A71800; MUID:99120557
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                                                            A; Cross-references:
                                                                                                  A; Molecule type: DNA
                                                                                                                       A; Status: preliminary
                                                                                                                                          A; Accession: A71890
                                                                                                                                                           A; Reference number: A71800;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Accession: A43511
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C; Accession: A46480;
                                                                                                                                                                                                                                                                                                                                                                                       RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-404 <OSM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Osman,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Status: preliminary;
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                                                                           1-259 <ARN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             similarity
9; Conser
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                                            :AE001508; GB:AE001439; NID:g4155338; PIDN:AAD06352.1; strain J99
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D.T.; King, B.L.; D.; Mills, S.D.;

Jiang, Brown,

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of the human gastric p

PID:9415

E.D.; Taylor,

Doig, P.C.;

Smith,

D.E.;

G.F

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Mismatches

Indels

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Gaps

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N;Alternate names: high affinity IgG receptor
C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C; Superfamily: Fc gamma rec
C; Keywords: immunoglobulin
F; 128-180/Domain: immunoglo
                                                                        F;127-179/Domain: immunoglobulin
                                                                                                               C; Superfamily: Fc gamma receptor
                                                                                                                              A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1;
                                                                                                                                                                                                                    A; Reference number: A43511;
                                                                                                                                                                                                                                       A; Title: Molecular cloning and
                                                                                                                                                                                                                                                                                             A; Note: sequence extracted
                                                                                                                                                                                                                                                                                                                                                                    A;Reference number: A46480;
A;Accession: A46480
                                                                                                                                                                                                                                                                                                                                                                                       A; Title: Structure and mapping of the gene encoding A; Reference number: A46480; MUID:92166399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fc gamma (IgG) receptor high affinity – mouse
                                                                                                                                                                                                                                                                                                                                                                                                                           Immunol. 148, 1570-1575,
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                                                                                                                                                                                                                                                         D.W.; Osman, N.;
ol. 144, 371-378,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GSLVTLNCETNL 230
                                                                                          immunoglobulin receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                         C.A.; McKenzie, I.F.; Hogarth, P.M.
                                                                                                                                                                                                                                                                          ; Tate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptor
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75.0%;
78.7%;
75.0%;
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                                                                                                                                                                                                                      and expression of MUID:90111035
                                                                                                                                                                                                                                                                                                                                                                                                                           1992
                                                                                                                                                                                                                                                                            NCBI backbone
B.; McKenzie,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              homology
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                                                                      homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2;
Score 48;
Pred. No.
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Pred. No.
                                                                                                               immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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DB 2
0.49;
                                                                                                                                                                                                                                       the mouse high affinity
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                                                                                                                                                                                                                                                                          (NCBIN:85205, NCBIN:85208, I.F.C.; Hogarth, P.M.
                 2
                                                                                                             homology
                 Length 404;
                                                                                                                                                                                                                                                                                                                                                                                                        mouse
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                                                                                                                                                                                                                                                                                                                                                                                                      high
                                                                                                                            PID:g200753
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                                                                                                                                                                                                                                         Fc receptor for
                                                                                                                                                                                                                                                                                                 NCBIN:85212,
                                                                                                                                                                                                                                                                                                                                                                                                          gamma
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submitted to the EMBL Data Library, A; Reference number: S33979
A; Accession: S33981
                                                                                                                                                                                                                                                                                                                                                                 vif protein - human immunodeficiency virus type 1
C;Species: human immunodeficiency virus type 1, HIV-1
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Accession: E64625
R;Tomb, J.F.; White, O.; Kerlavage, A.R.; Clayton, R.A.; Sutton, G.G.; Fleischmann, R.D. Peterson, S.; Loftus, B.; Richardson, D.; Dodson, R.; Khalak, H.G.; Glodek, A.; McKenne son, J.D.; Kelley, J.M.; Cotton, M.D.; Weidman, J.M.; Fujii, C.; Bowman, C.; Watthey, L. Nature 388, 539-547, 1997
Nature 388, 539-547, 1997
A;Authors: Wallin, E.; Hayes, W.S.; Borodovsky, M.; Karpk, P.D.; Smith, H.O.; Fraser, C. A;Title: The complete genome sequence of the gastric pathogen Helicobacter pylori.
A;Reference number: A64520; MUID:97394467
A;Accession: E64625
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Best Local Similarity
Thes 7; Conserv
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                                                                                                                                                                        A;Cross-references: EMBL:Z11530; C;Superfamily: AIDS vif protein
                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-192 <CAR>
A; Cross-references: EMB
                                                                                                                                                                                                                                                                                                                                                     C; Accession: S33981
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F;2-189/Domain: hydroxyethylthiazole kinase
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                                                                                                                                                                                                                                                                                                                                 R; Carlini,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hydroxyethylthiazole kinase (EC 2.7.1.50) – Helicobacter pylori (strain 26695)
C;Species: Helicobacter pylori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
                                                                                                                                                                                                                                                   Status: preliminary
                                                                                             Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Superfamily: hydroxyethylthiazole kinase; hydroxyethylthiazole kinase homology 
Keywords: phosphotransferase; thiamin biosynthesis 
:16-203/Domain: hydroxyethylthiazole kinase homology <HTK>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Status: preliminary; nucleic acid sequence not shown; translation not shown Molecule type: DNA Residues; 1-273 <TOM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pathway: thiamin biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Description: catalyzes the phosphorylation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cross-references: GB:AE000595; GB:AE000511; NID:g2313969; PIDN:AAD07891.1; PID:g23139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    120
126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           134 GSLVGISCESK 144
                                      1 GNLVTLSCE 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GNLVTLSCETK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GSLVGISCESK 130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GNLVTLSCETK 11
GNVVRLSCE
                                                                                               Similarity
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                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                             62.3%;
                                                                                                                                                                                          NID:g60192; PIDN:CAA77623.1; PID:g60195
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••
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 39;
Pred. No.
                                                                                             Score
Pred.
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Pred.
                                                                                                                                                                                                                                                                                                              November 1991
                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               39;
No.
                                                                                               ŏ.
                                                                                                               38;
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13;
                                                                                         DB
15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
14;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 by ATP of 4-methyl-5-(2-hydroxyethyl)-thia:
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                                                                                                               Length 192
                                                                           Indels
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                                                                         Gaps
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                                                                                                                                                                                            RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R; Carthew,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A36195
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hypothetical protein DKFZp
C;Species: Homo sapiens (m
C;Date: 11-Jun-1999 #seque
C;Accession: TO8678
C;Accession: TO8678
R;Mambutt, R.; Heubner, D
submitted to the Protein S
A;Reference number: Z16469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                developmental protein sina - fruit fly (Drosophila virilis)
C;Species: Drosophila virilis
C;Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 21-Feb-1997
C;Accession: A41544
                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Gene: FlyBase:sina
A;Cross-references: FlyBase:FBgn0003410
C;Keywords: DNA binding; nucleus; zinc finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cell 63, 561-577, 1990
A;Title: seven in absentia, a gene required for specification of R7 cell fate in the A;Reference number: A36195; MUID:91029488
A;Accession: A36195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    developmental protein sina - fruit fly (Drosophila C;Species: Drosophila melanogaster C;Date: 14-Dec-1990 #sequence_revision 13-Jan-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: FlyBase:FBgn0013142
C;Keywords: DNA binding; zinc finger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-314 <NEU>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Proc. Natl. Acad. Sci. U.S.A. 88, 10203-1
A; Title: Evolution of gene position: chro
A; Reference number: A41544; MUID:92052239
A; Molecule type: mRNA
A; Residues: 1-584 <WAM>
                                   A; Accession: T08678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 1-314 <CAR>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Cross-references:
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                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Status: preliminary
                                                                                                                                                                                                                                                                              90 GHLVCVSCRSKL 101
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                                                                                                                                                                             protein DKFZp564I1922.1 - human (fragment)
                                                                                               Heubner, D.;
                                                                                                                                      Carthew, R.W.; Rubin, G.M. Sci. U.S.A. 88, 10203-10207
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                                             GB:M38384; NID:g158466;
                                                                                                                                                        (man)
                                                                                                                                                                                                                                                                                                                                                                           58.3%;
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                                                                                               Mewes,
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Pred. No.
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Pred. No. 24;
                                                                            Database,
                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Drosophila melanogaster)
                                                                            Gassenhuber, J.; Wiemann,
se, May 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PID:g158467
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T-cell rece
C;Species:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Homo sapiens (man)
C;Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 30-May-1997
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A;Experimental source: fetal brain;
                                     RESULT
S38388
                                                                                                                                                                                                                                                                                                                                                                           R;Giegerich, G.; Pette, M.; Meinl, E.; Epplen, J.T.; Wekerle, H.; Hinkkanen, A. Eur. J. Immunol. 22, 753-758, 1992
A;Title: Diversity of T cell receptor alpha and beta chain genes expressed by human TA;Reference number: S57869; MUID:92192091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Species: Homo sapiens
C; Date: 27-Oct-1995 #sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: GB:M27388
C;Superfamily: immunoglobulin V
C;Keywords: T-cell receptor
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A; Status: preliminary
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A; Residues: 1-113 <KIM>
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                                                                                                                                                                                                                                                                     Molecule type: mRNA
Residues: 1-137 <GIE>
Superfamily: immunoglobulin
Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Accession: S57869
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Best Local
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 receptor beta chain precursor ies: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptor Mb11 beta chain - human
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8; Conser
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                                                                                                                                                                                                  60.7%;
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                                                                                                                                                                                                                                                                                       region;
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Pred.
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                      human (fragment)
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42;
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14;
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                                                                                                                                                                                                                   Length 137
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                            hypothetical protein slr1243 - Synechocystis sp. (strain PCC 6803)

C;Species: Synechocystis sp.

A;Variety: PCC 6803

A;Variety: PCC 6803

C;Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 20-Jun-2000

C;Date: 25-Apr-1997 #sequence_revision 25-Apr-1997 #text_change 20-Jun-2000

C;Accession: S74861

R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Asamizu, E.; Nakamura, Y.; I

R;Kaneko, T.; Sato, S.; Kotani, H.; Tanaka, A.; Wadda, T.; Watanabe, A.; Yamadi

O, K.; Okumura, S.; Shimpo, S.; Takeuchi, C.; Wadda, T.; Watanabe, A.; Yamadi

DNA Res. 3, 109-136, 1996
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     A;Reference number: $74322;
A;Accession: $74861
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-1064 <KAN>
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364

MUID:97061201

the unicellular cyanobacterium

Yamada,

; Miyajima, ada, M.; Yas Synechocys

2 NLVTLSCETKL 12

Conservative

54.5%;

Score 37; DB Pred. No. 43; 2; Mismatches

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Length 382

0;

Gaps

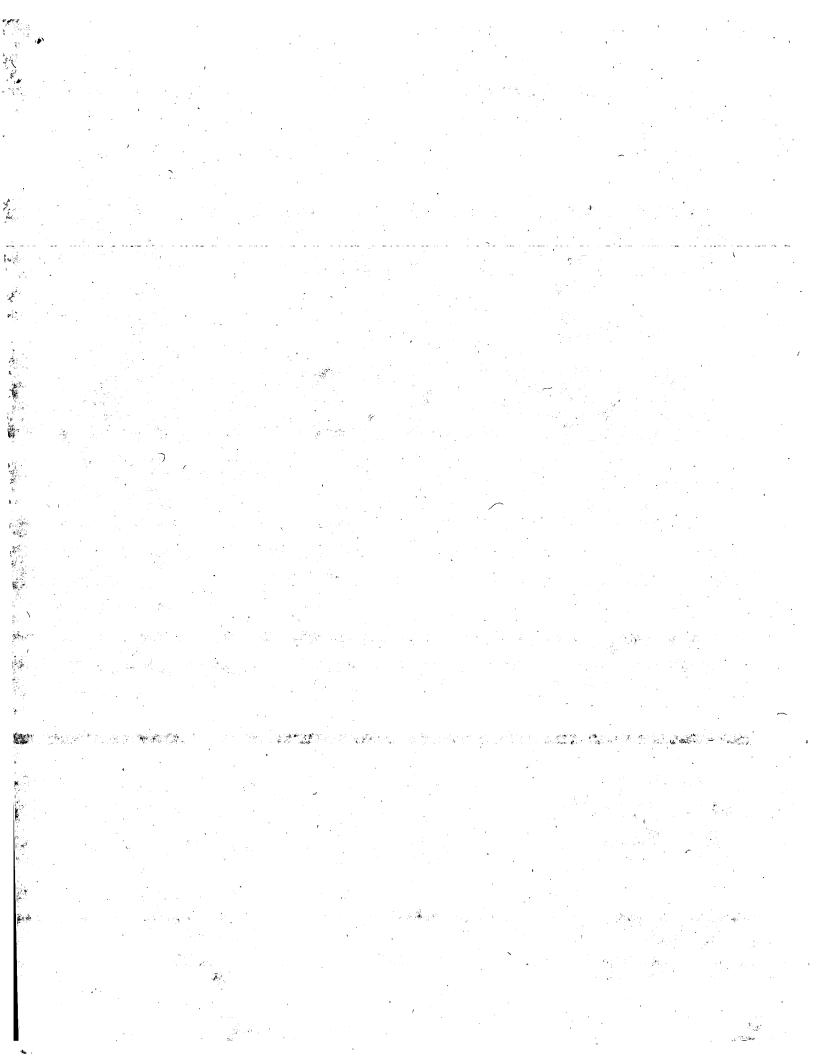
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C; Superfamily: immunogiobut
C; Keywords: T-cell receptor
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    A; Gene: MTH141
C; Superfamily:
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                                                                   A;Cross-references: GB:AE000903; GB:AE000666; NID:g2622514; A;Experimental source: strain Delta H
                                                                                                                                                                                                                                                                                                                                                                                probable cell division control protein 6 Methanobacterium thermoautotrophicum (stra C; Species: Methanobacterium thermoautotrophicum
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                                                                                                               A; Molecule type: DNA
A; Residues: 1-382 <MTH>
                                                                                                                                                                            A; Title: Complete genome sequence of Methanobacterium thermoautotrophicum Delta H: A; Reference number: A69000; MUID:98037514
A; Accession: B69055
                                                                                                                                                                                                                                                                                                                   R; Smith, D.R.;
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79, 7135-7155, 1997
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A;Cross-references: EMBL:D90909; GB:AB001339; NID:g1652844; PIDN:BAA17822.1; PID:g165290 A;Note: the nucleotide sequence was submitted to the EMBL Data Library, June 1996 C;Superfamily: Synechocystis hypothetical protein slr1243
                                                                                                                                Query Match 60.7%; Score 37; DB 2; Length 1064; Best Local Similarity 45.5%; Pred. No. 1.1e+02; Matches 5; Conservative 4; Mismatches 2; Indels
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P29304 drosophila
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	Q99795	014953	P43631	Q9x5z9	031213	Q10343	P54356	Q62230	290703	015394	P27462	P04217
ı	homo sapien	homo sapien	homo sapien	streptomyce	streptomyce	schizosacch	drosophila	mus musculu	gallus gall	homo sapien	broad bean	homo sapien

## ALIGNMENTS

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                                                                        "Structure and mapping of the gene en gamma RI and chromosomal location of J. Immunol. 148:1570-1575(1992).
                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vert
Mammalia; Eutheria; Rodentia; Sciurognathi;
  -1- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLO)
AFFINITY RECEPTOR.
-1- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN
-1- TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
-1- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2
                                                                                                                                                            MEDLINE-92166399; PubMed=1531670;
Osman N., Kozak C.A., McKenzie I.F.,
                                                                                                                                                                                                                                                            MEDIINE=90111035; PubMe Sears D.W., Osman N., 7 "Molecular cloning and receptor for IgG.";
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IMMUNOSLOBULIN GAMMA FC RECEP
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d expression
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2C2AA8103ECF16E6 CRC64;
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ion of the
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                                                                                                                F., Hogarth P.M.;
e encoding mouse
of the human Fc
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                                                                               OF IMMUNOGLOBULINS
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thi; Muridae;
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e mouse high affinity
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; Murinae; Mus
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RESULT 3
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.Best Local S
Matches
gastric pathogen Helicobacter pylori.";
Nature 397:176-180(1999).
-!- CATALYTIC ACTIVITY: ATP + 4-METHYL-5-(2-HYDROXYETHYL)-THIAZOLE
ADP + 4-METHYL-5-(2-PHOSPHOETHYL)-THIAZOLE.
-!- COFACTOR: MAGNESIUM (BY SIMILAPTHY)
                                                                                                                       SEQUENCE FROM N.A.
MEDLINE-99120557; PubMed=9923682;
Alm R.A., Ling L.-S.L., Moir D.T.,
Smith D.R., Noonan B., Guild B.C.,
Tummino P.J., Caruso A., Uria-Nicke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CARBOHYD
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01-OCT-2000
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DOMAIN
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PIR; A43511; A43511.
PIR; A46480; A46480.
                                                                                                 Irust T.J.;
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InterPro; IPR003006; -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00047; ig;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or send an
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9; Conser
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Last annotation update)
KINASE (EC 2.7.1.50) (4-
KINASE) (THZ KINASE) (TH
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                                                                                                                                                                                                                                                 (Campylobacter pylori ; epsilon subdivision;
                                                                                                                       Moir D.T., King B.L., Brown E.D., Doig uild B.C., deJonge B.L., Carmel G., Uria-Nickelsen M., Mills D.M., Ives C.
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(TH KINASE).
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RESULT 4
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Cotton M.D., Weldman J.M., Fujli C., Bowman C., Watthey L., Wa
Hayes W.S., Borodovsky M., Karp P.D., Smith H.O., Fraser C.M.,
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01-OCT-2000 (Rel. 40,
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HYDROXYETHYLTHIAZOLE)
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                                                              between the Swiss Institute of Bioinformatics are the European Bioinformatics Institute. There are
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MEDLINE=97394467; Pu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                            Nature 388:539-547(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=210;
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SIMILARITY: BELONGS TO THE THZ KINASE FAMILY.
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                   non-profit instituend this statement
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4-METHYL-5-(2-PHOSPHOETHYL)-THIAZOLE
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                                            Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J., Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C., Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
                                                                                                                                                                                                                                                                                                              George R.A., Lewis S.B., Natural D.D., Zhang Q., Chen L.X., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Nelson C.R., Miklos G.L.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. MEDLINE-91029488; PubMed-2146028; Carthew R.W., Rubin G.M.;
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                                                                                                                                                                  Durbin K.J.,
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Amanatides P.G., Scherer S.E., Li P.W.,
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                                                                                                                                                                , Delcher A., Deng Z., Mays A.D., Dew T., Dietz S.M., oup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn Evangelista C.C., Ferraz C., Ferriera S., Fleischmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Drosophilidae;
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188
27514 MW;
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Reinert K.,

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Nixon K., Nusskern D.R., Pacle ., 'Pan S., Pollard J., Puri V.,

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Best Local
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01-DEC-1992 (Rel. 24, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation updat
DEVELOPMENTAL PROTEIN SEVEN IN ABSENTIA.
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P29304;
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                                                                                            MEDLINE=92052239; PubMed=1946441;
Neufeld T.P., Carthew R.W., Rubin G.M.;
"Evolution of gene position: chromosomal arrangement and
comparison of the Drosophila melanogaster and Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Science 287:2185-2195(2000).
-I- FUNCTION: REQUIRED FOR SPECIFICATION OF R7 PHOTORECEPTOR CELL FARE IN THE DROSOPHILA EYE. POSSIBLY ACTS BY REGULATING GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J. S., Zhan M., Zhang G., Zhao Q., Zheng L., Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu X., Smith H.O., Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.; "The genome sequence of Drosophila melanogaster.";
                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Arthropoda; Tracheata; Pterygota; Neoptera; Endopterygota; Diptera
Ephydroidea; Drosophilidae; Drosophila.
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entities requires a license agreement (See
or send an email to license@isb-sib.ch).
                                                         Proc.
                                                                                and Rh4
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Eukaryota; Metazoa;
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the European Bioinformatics Institute
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
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c. Natl. Acad. Sci. U.S.A. 88:10203-10207(1991).
FUNCTION: REQUIRED FOR SPECIFICATION OF R7 PHOTORECEPTOR CELL
FATE IN THE DROSOPHILA EYE. POSSIBLY ACTS BY REGULATING GENE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 314 AA;
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RESULT 7
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Best Local
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ZN_FING
                                                      transactivator and class II MHC.";
Cancer Immunol. Immunother. 42:99-107(1996).
Cincer Immunol. Immunother. 42:99-107(1996).
II PROMOTER: ACTIVATION IS VIA THE PROXIMAL ACTIVITY OF THE HLA CLII PROMOTER; ACTIVATION IS VIA THE PROXIMAL PROMOTER. MAY ACT IN BINDING OF IN VITRO TRANSLATED CIITA WAS DETECTED. MAY ACT IN COACTIVATOR-LIKE FASHION THROUGH PROTEIN-PROTEIN INTERACTIONS
                                                                                                                                                                                                                                                                                             "Mouse class II transactivator: cDNA sequence and am comparison with the human class II transactivator."; Immunogenetics 45:220-222(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  P79621; Q31115;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
MHC CLASS II TRANSACTIVATOR CIITA.
MHC2TA OR CIITA OR C2TA.
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MEDLINE=9618886; PubMed-862527;
Panelli M.C., Wang E., Shen S., S
Hersh E.M., Stopeck A., Gangavall
                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
Eukaryota; Metazoa; (
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                                                                                                                                                                poorly responsive to IFNgamma
                                                                                                                                                                                         Akporiaye E.T.
                                                                                                                                                                                                                                                                                                                                                                    STRAIN-NOD;
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DOMAIN
                                                                                                                                       mmunogenicity is accompanied by induction
                                                                                                                                                                             'Interferon gamma (IFNgamma) gene transfer of an EMT6 tumor that
                                                                                                                                                                                                                                                                   SEQUENCE OF 878-1020 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90 GHLVCVSCRSKL
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TISSUE SPECIFICITY: IN MANY OMMATIDIAL PRECURSOR CELLS.
CONTACTING FACTORS BINDING TO THE PROXIMAL MHC CLASS II PROMOTER, TO ELEMENTS OF THE TRANSCRIPTION MACHIMERY, OR BOTH. ALTERNATIVEL IT MAY ACTIVATE HLA CLASS II TRANSCRIPTION BY MODIFYING PROTEINS THAT BIND TO THE MHC CLASS II PROMOTER (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SWISS-PROT entry is copyright. It is produced through a
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k A., Gangavalli
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TO DICTYOSTELLIUM DG17 (AA 25-67).
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lli R., Barber
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                                                                                                                                                increase in tumor of a mouse class
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                               use by non-profit institutions as long modified and this statement is not removed.
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                                                                                                                                                                                                                                                                                                                            entities requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                  EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE-Bone marrow;
MEDLINE-97191544; PubMed-9039502;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RW1 PROTEIN
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                                                                                                  Similarity 7; Conser
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Pred. No.
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PRECURSOR (T-CELL

SURFACE ANTIGEN

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CD4_SALSC
ID CD4_SALSC
AC 029037;
DT 01-NOV-1997
DT 01-NOV-1997
DT 15-JUL-1999
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IL18_RJ
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ul-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation updat
Ţ-CELL SURFACE GLYCOPROTEIN CD4 pproviscon "
                                                                                                                                                                                                                                                                     CONFLICT
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
10-OCT-2000 (Rel. 40, Last annotation update)
INTERLEUKIN-18 PRECURSOR (IL-18) (INTERFERON-GAMMA INDUCING FACTOR)
(IFN-GAMMA-INDUCING FACTOR) (INTERFERON-I GAMMA) (IL-1 GAMMA).
                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by a
entitles requires a license agreement (See http://www.isb-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-SPRAGUE-DAWLEY; TISSUE-Adrenal gland; MEDLINE-97152963; PubMed-8999896; Contl B., Jahng J.W., Tintl C., Son J.H., Jo "Induction of interferon-gamma inducing fact
                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                  EMBL;
                                                                                                                                                                                                                                                                                                                                                             EMBL;
                                                                                                                                                                                                                                                                                                                                                                            EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                      or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PubMed=9702748;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-SPRAGUE-DAWLEY; TISSUE-Brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cortex.";
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IL18_RAT
                                                                                                                                     105
                                                                                                                                                                                                                                                                                              VARSPLIC
                                                                                                                                                                                                                                                                                                            CHAIN
                                                                                                                                                                                                                                                                                                                                      Cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MO1. PSych. 3:362-366(1998).
-i- FUNCTION: AUGMENTS NATURAL KILLER CELL ACTIVITY IN SPLEEN CELLS
AND STIMULATES INTERFERON GAMMA PRODUCTION IN THELPER TYPE I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Culhane A.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rattus norvegicus (Rat).
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    Biol. Chem.

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                                                                                                                                                                                                    Local
                                                                                                                                                 1 GNLVTLSCETKL
| : ||||: |:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBCELLULAR LOCATION: SECRETED. SIMILARITY: BELONGS TO THE IL-18 FAMILY.
                                                                                                                                     GRMSTLSCKNKI 116
                                                                                                                                                                                                                                                                                                                                                 077777; AAC53010.1; -
AJ222813; CAA11001.1;
                                                                                                                                                                                                                                                                                                                                                                            U77776;
                                                                                                                                                                                      l Similarity
6; Conserv
                                                                                                                                                                                                                                                                                                                    Alternative splicing.

36 BY SIMILARITY
                                                                                                                                                                                                                                                      37
121
4
48
194 AA;
                                                                                                                                                                                        Conservative
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                                                                     STANDARD;
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139
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Rodentia;
                                                                                                                                                                                                   59.0%;
                                                                                                                                                                                                                                                        22303 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       interleukin-18 cDNA.";
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                                                                                                                                                                                                   Score 36;
Pred. No.
                                                                                                                                                                                                                                                      MISSING (IN ISOFORM ALPHA).
MS -> IP (IN REF. 2).
I -> M (IN REF. 2).
; E2089AD6F1798450 CRC64;
                                                                                                                                                                                                                                                                                                          INTERLEUKIN-18
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                 update)
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RESULT 11
VPL_HAEIN
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Query Match
Best Local
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01-NOV-1995
01-OCT-2000
                                                             VPL_HAEIN
P44233;
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SIGNAL
Haemophilus influenzae
Bacteria; Proteobacter
                            MU-LIKE PROPHAGE
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PRINTS; PR00692; CD4TC,
PRINTS; PR00692; CD4TC,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    entities requires a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Ve
Mammalia; Eutheria; Primates; Platyrrhini;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Saimiri sciureus
                                                                                                                                                                                                                                                                                                                                    DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T4/LEU-3).
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                                                                                                                  34 GEVVELPCETSL
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                                                                                                                                 GNLVTLSCETKL
                                                                                                                                                     Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                             Lipoprotein;
Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                              IPR000973; -.
                                    (Rel.
                                                                                                                                                                                                 457
                                                                                                                                                    Conservative
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                                                                       STANDARD;
                            FLUMU
                                    32, Created)
32, Last sequence 40, Last anno
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417
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                          Last sequence update)
Last annotation update)
U TAIL SHEATH PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                    Transmembrane;
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gamma
                                                                                                                                                            Score 36;
Pred. No.
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BY SIMILAL
BY SIMILAL
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
                                                                                                                                                     Pred. No. 20;
; Mismatches
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CYTOPLASMIC (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                  EXTRACELLULAR
                                                                       PRT;
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subdivision;
                                                                                                                                                                                                                         SIMILARITY
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D (GLCNAC. . .)
D (GLCNAC. . .)
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ni; Cebidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   as
                                                                                                                                                                                                                                                                                                                                  (POTENTIAL)
Pasteurellaceae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  its content
                                                                                                                                                                     Length 457;
                                                                                                                                                                                                                                                  ) (POTENTIAL).
) (POTENTIAL).
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Cebinae; Saimiri.
                                                                                                                                                                                                                                                                                                                                                                    T-cell; MHC;
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RESULT
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Best Local S
Matches
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                                                                                                                                                                                                                                                                                                                                                                                               DROME
                                                                                                                                                                                                                     AHCY89E OR PH200 OR CG8956.

Drosophila melanogaster (Fruit fly).

Eukaryota; Metazoa; Arthropoda; Tracheata; Herterygota; Neoptera; Endopterygota; Diptera; Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fleischmann R.D., Adams M.D., White O., Clayton R.A., Kirkness E.F., Kerlavage A.R., Bult C.J., Tomb J.F., Dougherty B.A., Merrick J.M., McKenney K., Sutton G., Fitzhugh W., Fields C.A., Gocayne J.D., Scott J.D., Shirley R., Liu L.-I., Glodek A., Kelley J.M., Weidman J.F., Phillips C.A., Sprigs T., Hedblom E., Cotton M.D., Utterback T.R., Hanna M.C., Nguyen D.T., Saudek D.M., Brandon R.C., Fine L.D., Fritchman J.L., Fuhrmann J.L., Geoghagen N.S.M., Gnehm C.L., McDonald L.A., Small K.V., Fraser C.M., Smith H.O.,
MEDLINE=95396803; PubMed=7667301;
Martin C.H., Mayeda C.A., Davis C
Mathog D.R., Celniker S.E., Lewis
                                            SEQUENCE FROM N.A. STRAIN=CANTON-S;
                                                                                     trans-regulatory homeoprotein.";
EMBO J. 7:3223-3231(1988).
                                                                                                         MEDLINE-89030618; PubMed=2903049;
Delorenzi M., Ali N., Saari G., Henry C., Wilcox M., Bienz M.;
Pevidence that the Abdominal-B r element function is conferred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed
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                                                                                                                                                            SEQUENCE FROM N.A.
TISSUE=Embryo;
                                                                                                                                                                                                                                                                                                                                                                                   SAH2_DROME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hypothetical SEQUENCE 41
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NCBI_TaxID=727;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Whole-genome random sequencing and assembly of Haemophilus influenzae Rd.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Venter J.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                           163
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 54.96; Conservative
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487 AA; 53143
                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
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54.5%;
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   Lewis
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                   PRT;
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MU PROTEIN L.
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               Ericsson C.L.,
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   Palazzolo M.J.;
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a; Brachycera;
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Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,

Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,

A Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,

Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,

A Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Fleischmann W.,

A Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.,

A Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,

A Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,

A Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,

A Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,

A Harris N.L., Harvey D., Heiman T.J., Wei M.-H., Ibeywam C.,

A Harris N.L., Harvey D., Heiman T.J., Wei M.-H., Ibeywam C.,

A Harris N.L., Harvey D., Heiman T.J., Wei M.-H., Ibeywam C.,

A Harris N.L., Harvey D., Heiman T.J., Wei M.-H., Ibeywam C.,

A Harris N.L., Kalush F., Karpen G.H., Ke Z., Kulp D., Lai Z.,

A Kimmel B.E., Kodira C.D., Kraft C., McLeod M.P., McPherson D.,

A Merkully G., Milskins N. W. McCharry C., McCherle I. McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherle I. McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherle I. McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherry C.,

A Markully G., Milskins N. W. McCharry C., McCherry C.,

A Markully G., Milskins N. McCharry C., McCherry C.,

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A Markully G., Milskins N. M. McCharry C.,

A Markully G
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Sutton G.G.,
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George R.A., Lewis S.E., Richards S., Ashburner M.,
                                                                                                                                                                                                                    Pfam; PF00670;
                                                                                                                                                                                                                                                                                                                        EMBL; U31961; AAA84400.1; -.
EMBL; AE003715; AAF55367.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified and this statement
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                                                                                                                                           PROSITE; PS00738; ADOHCYASE_1; PROSITE; PS00739; ADOHCYASE_2;
                                                                                                                                                                                                                                                                                      FlyBase; FBgn0015011; Ahcy89E
                                                                                                                                                                                                                                                                                                                                                                                             EMBL; X13168; CAA31566.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or send an email to license@isb-sib.ch).
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                                                                                                          Hydrolase;
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                                                                                                                                                                                                                                                         [nterPro;
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SIMILARITY: BELONGS TO THE ADENOSYLHOMOCYSTEINASE FAMILY
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TISSUE SPECIFICITY: EXPRESSED IN EXTENDI
SOMATIC MESODERM, YOLK CELLS AND MIDGUT
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NAD;
275
127
484
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                                                                                                              One-carbon
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                license agreement (See http://www.isb-sib
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                                                                                                              metabolism
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NAD (POTENTIAL).
T -> A (IN REF. 1).
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Zhu X., Smith H.O.
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070472;
01-0CT-2000
01-0CT-2000
01-0CT-2000
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                                                                                                     P13944: Q04509;
01-JAN-1990 (Rel. 13, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
COLLAGEN ALPHA 1(XII) CHAIN PRECURSOR (FIBROCHIMERIN).
                                                                                                                                                                                                                              1131
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                                                                                                                                                                              CHICK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                   Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                                                                                                                                                             use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wilkinson R., Fitter S., Tscharke D., Simmons A.;
Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: MAY PLAY A ROLE IN THE IMMUNE RESPONSE TO VIRAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090
                                                        Archosauria; Aves;
                                                                                              COL12A1
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                                                                                                                                                                                                                                                                                                                                                                                   or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                between the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There a
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                                                                     Craniata; Vertebrata; Euteleostomi;
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"Large and small splice variants of collagen XII: differential expression and ligand binding."; |
J. Cell Baol. 130:1005-1014(1995).
-1- FUNCTION: TYPE XII COLLAGEN INTERACTS WITH TYPE I COLLAGEN-
CONTAINING FIBRILS, THE COLL DOMAIN COULD BE ASSOCIATED WITH THE
SURFACE OF THE FIBRILS, AND THE COL2 AND NC3 DOMAINS MAY BE
LOCALIZED IN THE PERIFIERILLAR MATRIX.
-1- SUBUNIT: TRIMER OF IDENTICAL CHAINS EACH CONTAINING 190 KDA OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Trueb J., Trueb B.;
"The two splice variants of collagen XII share a Biochim. Biophys. Acta 1171:97-98(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Type XII collagen: distinct extracellular matrix component discovered by cDNA cloning."; proc. Natl. Acad. Sci. U.S.A. 84:6040-6044(1987).
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Gordon M.K., Gerecke D.R., Dublet B., va
"Type XII collagen. A large multidomain
homology to type IX collagen.";
J. Biol. Chem. 264:19772-19778(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "The complete primary structure of type XII collagen shows a chimeric molecule with reiterated fibronectin type III motifs, von Willebrand factor A motifs, a domain homologous to a noncollagenous region of type IX collagen, and short collagenous domains with an Arg-Gly-Asp
                                                                    modified and this statement is not removed.
                                                                                                                          the European
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Embryo;
MEDLINE=93042014; PubMed=1420368;
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Nishida Y., Obara M.,
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                                                                                                                                                                                                                                            PTM: GLYCOSYLATED, CHONDROITIN-SULFATED (BY SIMILARITY: CONTAINS 4 WAFA DOMAINS.
SIMILARITY: CONTAINS 18 FIBRONECTIN TYPE III-LIKE DOMAINS.
SIMILARITY: BELONGS TO THE FAMILY OF FIBRIL-ASSOCIATED COLLAGENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LIGAMENTS, PERICHONDRIUM, AND PERIOSTEUM, ALL DENSE CONNECT TISSUES CONTAINING TYPE I COLLAGEN.

DOMAIN: THIS SEQUENCE DEFINES FIVE DISTINCT DOMAINS, TWO TR. HELICAL DOMAINS (COL1 AND COL2) AND THREE NONTRIPLE-HELICAL DOMAINS (COL1 AND NC3).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NONTRIPLE-HELICAL SEQUENCES.

ALTERNATIVE PRODUCTS: 2 ISOFORMS; A LONG FORM (SHOWN HERE) AND A SHORT FORM; ARE PRODUCED BY-ALTERNATIVE SPLICING. THE FINAL TISSUI FORM OF COLLAGEN XII MAY CONTAIN HOMOTRIMERS OF EITHER THE LONGER OR THE SHORTER ISOFORM OR ANY COMBINATION OF LONG AND SHORT ISOFORM CHAINS. ONLY THE LONG VARIANT IS A PROTEOGLYCAN. THE LARGI ISOFORM HAS MORE RESTRICTED EXPRESSION IN EMBRYONIC TISSUE THAN
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                                                                                                                                              SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN
requires a license agreement (See http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
                                                                                                                                                                                                                                    INTERRUPTED HELICES (FACIT).
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Bioinformatics Institute.
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)bara M., Kimata K.;
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                                                                                        rmatics Institute. There are no restinstitutions as long as its content
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                                                           NONHELICAL REGION (NC1).

ASP/GLU-RICH (ACIDIC).

ARG/LYS-RICH (BASIC).

TO CHONDROITIN SULFATE (POTENTIAL).

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TO CHONDROITIN SULFATE (POTENTIAL).

CELL ATTACHMENT SITE (POTENTIAL).

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MEDLINB-88166903; PubMed=3127222;

Brueggemann M., Delmastro-Galfre P., Waldmann H., Calabi F.;

"Sequence of a rat immunoglobulin gamma 2c heavy chain constant

"Sequence of a rat immunoglobulin gamma 3.";

region cDNA: extensive homology to mouse gamma 3.";

Eur. J. Immunol. 18:317-319(1988).
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P20762;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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01-FEB-1991 (Rel. 17, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
IG GAMMA-2C CHAIN C REGION.
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V -> F (IN REF. 2).
QP -> AG (IN REF. 3)
B MW; 094285AFE7F346CF
                                                       Score 35; DB 1;
Pred. No. 22;
1; Mismatches
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Pred. No. 1.4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     329 AA.
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Search completed: June Job time: 564 sec 4, 2001, 12:24:02

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Minimum DB
Maximum DB
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Match
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13.062 Million cell updates/sec
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sp_virus:*
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Q9JDG0
Q9JDE7
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O78109 mus musculu
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01-FEB-1997 (TrEMBLrel.
01-FEB-1997 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
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# ALIGNMENTS

02, 15,

Created)

PRT;

374 AA

Last sequence update)
Last annotation update)

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RESULT 2
Q92495
ID Q92495
                                                                                                                                                 Query Match
Best Local Similarity
Matches 12; Conserv
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                               PFAM; PF00047; ig; 3. PRODOM; PD002534; -;
                                                                                                                                                                                                                                                                                                   EMBL; L03418; AAA36049.1; HSSP; P12319; FALT.
                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-93055454; PubMed=1430234; Porges A.J., Redecha P.B., Doebel Kimberly R.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                 205 GNLVTLSCETKL 216
                                                                                                    1 GNLVTLSCETKL 12
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                                                                                                                                                                                                                                                                                                                                                                 Fc gamma receptor I family gene products in human mononuclear
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   PRELIMINARY;
                                                                                                                                                    Conservative
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42632 MW;
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                                                                                                                                                 Score 61; DB 4;
Pred. No. 0.0003
); Mismatches
   PRT;
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   375 AA.
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                                                                                                                                                                                 Length 374;
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7. Exp. ...
79155
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                                                  high
                                                                                            Yan
                                                                                                                                            Bovidae; Bovinae; NCBI_TaxID=9913;
                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pe
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                                                                                                           SEQUENCE FROM N.A.
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Eukaryota; Metazoa; Chordata;
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01-FEB-1997 (TrEMBLrel.
01-FEB-1997 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
Receptor.
                 EMBL; AF162866; AAF80477.1;
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                                (an Y. Li X., Wang A., Zhang G.;
Molecular cloning and identification of full-length cDNA encoding
ligh affinity Fc receptor for bovine IgG (Fc gamma RI).";
Pet. Immunol. Immunopathol. 75:151-159(2000).
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C GAMMA RIB-FC GAMMA RECEPTOR).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A84D464C70DD0F91 CRC64;
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RESULT
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Matches 11; Conser
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Best Local :
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Q9Y1V6;
Q1-NOV-1999
Q1-NOV-1999
Q1-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                      "Two localization pathways of maternal RNAs at the post-cytoplasm in early ascidian embryos."; Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases. EMBL; ABO2935; BAA81908.1; -
                                                                                                                                                                                                                       Q38131;
Q38131;
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PROSITE; I
                                                                                                                                                                                01-NOV-1996 (TrEMBLrel.
01-NOV-1996 (TrEMBLrel.
01-NOV-1998 (TrEMBLrel.
STRUCTURAL PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Halocynthia roretzi (Sea squirt).
Eukaryota; Metazoa; Chordata; Urochordata;
Stolidobranchia; Pyuridae; Halocynthia.
       Mol. Microbiol. 19:1343-1355(1996).
                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                              MEDLINE=96332669;
van Sinderen D., 1
                                                                 SEQUENCE FROM N.A.
                                                                                                               MEDLINE=96332668; PubMed=8730874; Nauta A., van Sinderen D., Karsen
                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                               Viruses.
                                                                                                                                                                       Bacteriophage rlt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HRPET-3
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                           Venema G., Nauta A.;
"Sequence analysis and molecular characterization
                                                                                             isolated
                                                                                                                                                    NCBI_TaxID=43685;
                                                                                                       "Inducible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=7729;
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3; PS01179; PID; 1.
690 AA; 75525 P
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                                                                                  Sinderen D., Karsens H., Smit E., Venema G., K
ne expression mediated by a repressor-operator
Lactococcus lactis bacteriophage rlt.";
1. 19:1331-1341(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA;
                                               Karsens
                                              PubMed=8730875;
Karsens H., Kok
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Pred. No. 0.57
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Query Match Best Local S Matches 8

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65.6%;

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Q9U6A9;
01-MAY-2000
01-MAY-2000
01-JUN-2000
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Q97753;
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                          Park J.-H., Jensen B.C., Pars
"A novel nucleolar G-protein
Submitted (SEP-1999) to the I
EMBL; AF189284; AAF01061.1;
                                                                                                                   NOG1.
                                                                                                                                                                                                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae;
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                                                             STRAIN-ISTAR1;
                                                                                                 Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                     "HIV-1 env sequence variation in brain related neurologic disease.";
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                                                                                      Pukaryota; Euglenozoa;
VCBI_TaxID=5691;
                                                                                                         rypanosoma
                                                                     SEQUENCE FROM N.A.
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                  IPR000765;
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GTP10BG.
A; 74797 MW;
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72.7%;
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3244326657BDA3E6 CRC64;
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                                                                                                Trypanosomatidae;
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SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
Amano H., Kato K., Miyamura T.
Amano H., Kato K., Miyamura T.
Sequence analysis of Yaba vir
Submitted (MAR-1999) to the EV
Submitted (MAR-1999) to the EV
EMBL; AB025319; BAA88969 1; -
                                                                                                                                                    Q9QAY4;
01-MAY-2000 (
01-MAY-2000 (
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YB-E11R.
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01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=COLUMBIA;
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itted (JAN-1999)
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AB022215; BAB01780.1; -
NCE 428 AA; 50007 MW;
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Similarity - 71.4%;
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62.3%;
54.5%;
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                                                                 Yaba virus DNA.";
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dicots; Rosidae;
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078242;
01-NOV-1996
01-NOV-1996
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                                                                                                                                                                                               Q9V539
Q9V539;
                                                                                                                                                                                                                                                                                                                                                               PRINTS; PROUVER 192 AA; 22543 MW;
                                                                                  Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Inse
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Ephydroidea; Drosophilidae; Drosophila.
MEDLINE=20196006; PubMed=10731132;
Adams M.D., Celniker S.E., Holt R.A.,
Amanatides P.G., Scherer S.E., Li P.W
                                                                                                                                     BCDNA: GH08789
                                                                                                                                               01-MAY-2000 (TremBLrel.
01-MAY-2000 (TremBLrel.
01-OCT-2000 (TremBLrel.
BCDNA:GH08789 PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
                                      STRAIN-BERKELEY
                                                 SEQUENCE FROM N.A.
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Borsetti A., Saggio I., Verani P., Ros
Int. J. Immunopharmacol. 3:17-23(1990)
EMBL, Z11530; CAA77623.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               clones from
                                                                         NCBI_TaxID=7227
                                                                                                                                                                                                                                                                                                                                                                                                                   INTERPRO; IPR000475; PFAM; PF00559; Vif; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Federico M., Titti F., Butto S., Orecchia A., Car
Macchi B., Mangiano N., Verani P., Rossi G.,
"Biologic and molecular characterization of produ
clones from HUT-78 cells infected with a patient
AIDS Res. Hum. Retroviruses 5:385-396(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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                                                                                                                                                                                                            PRELIMINARY;
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                                                                                                                                                                                                          PRT;
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                                                                                                          Tracheata; Hexapoda; Insecta;
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P.W.,
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Evans C.A., Gocayne J.D., ...
, Hoskins R.A., Galle R.F.,
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                                                                                                                                                                                                                                                                                                                                           Length 192;
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isolate.";
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Homo sapiens (Human).

Homo sapiens (Human).

Homo sapiens (Human).

Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Zhu Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.; "The genome sequence of Drosophila melanogaster.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Shue B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T., Spier E., Spradling A.C., Stapleton M., Strong R., Sun E., Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X., Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng
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Beeson K.Y.,
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PFAM; PF02141;
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EMBL; AE003834; AAF58981.1; -
FLYBASE; FBgn0027561; BcDNA:GH08789
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Sutton G.G.,
                                                                             TISSUE=BRAIN;
                                                                                               SEQUENCE FROM N.A.
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435 AA; 497
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Evangelista C.C., Ferraz C.
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50.0%;
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Pred. No.
                                                                                                                                               Craniata; Vertebrata; Catarrhini; Hominidae
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3., Helt G., Nelson C.
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15;
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X., Zhu S., Zhu X.,
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Baldwin D.,
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Wambutt R., Heubner D., Mewer Submitted (MAY-1999) to the EMBL; AL049946; CAB43220.1; HSSP; P56276; lTLK.

Mewes H.W., Gassenhuber J., Wiemann the EMBL/GenBank/DDBJ databases.

INTERPRO; IPR003006; -

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PF00047; 1g;

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078109;
01-NOV-1998 (TrEMBLrel. 08, C
01-NOV-1998 (TrEMBLrel. 08, L
01-JUN-2000 (TrEMBLrel. 18, L
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NON_TER
SEQUENCE 58
Q9TPP1;
01-MAY-2000 (
01-MAY-2000 (
01-MAY-2000 (
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (OCT-1998) to the EMBL; AF042159; AAC34367.1; EMBL; AF100710; AAF06839.1;
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Mus musculus (Mouse).
Eukaryota, Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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117 GDVVTVACEAK 127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EQUENCE FROM N.A.
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                                                                                                                                                                                                                                           GNLVTLSCET
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6; Conserv
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8; Conserv
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(TrEMBLrel.) (TrEMBLrel.)
                                                                                                                                                                                                                                                                                                                                                                                                 1054 AA;
                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the Major Histocompatibility Complex ous forms of the mouse transactivator
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CIITA FORM IV.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9C75943922BADCA9 CRC64;
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r Histocompatibility Complex
of the mouse transactivator
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36;
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                                                                                                                                                                                                                                                                                                                              Length 1054;
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Search completed: June Job time: 585 sec

4

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RESULT
Q9NR99
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Best Local Similarity
Watches 8; Conserv:
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                                                                                          Query Match
Best Local Similarity 54.
Matches 6; Conservative
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Q9NR99;
01-OCT-2000
01-OCT-2000
2361
                                                                                                                                                                                                       Crowl R.M., Luk D.;

'Identification of the gene encoding Adlican, a novel prexpressed in human arthritic tissues.";

Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AF245505; AAF86402.1;

EMBL; AF245505 AAF86402.1;

SEQUENCE 2828 AA; 312292 MW; A18377D8554F1FE1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-97327562; PubMed-9184229; Muhlethaler-Mottet A., Otten L.A., Steimle V., Mach B.; Muhlethaler mottet a., Otten L.A., Steimle V., Mach B.; Expression of MHC class II molecules in different cellular functional compartments is controlled by differential usage multiple promoters of the transactivator CITTA."; EMBO J. 16:2851-2860(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases. EMBL; AF100709; AAF06838.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Otten L.A., Steimle V., Bontron S., Mach B.; "Equivalent expression of the Major Histocompatibility Complex genes through the various forms of the mouse transactivator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-BALB/C
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                                                                                                                                                                                                                                                                                                                                                                        TISSUE=PLACENTA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADLICAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mammalia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
                                              1 GNLVTLSCETK
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GDVVTVACEAK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) (TrEMBLrel.) (TrEMBLrel.) (TrEMBLrel.
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15,
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Last sequence update)
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                                                                                               Score 38; DB Pred. No. 97; 4; Mismatches
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
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40;
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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence:
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                                                                                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                      Score
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Gapop 10.0 , Gapext 0.5
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61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Geneseq_0401:*
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/SIDS6/gcgdata/geneseq/geneseqp/AA1987.DAT:
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/SIDS6/gcgdata/geneseq/geneseqp/AA2001 .DAT:
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/SIDS6/gcgdata/geneseq/geneseqp/AA2000.DAT:*
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/SIDS6/gcgdata/geneseq/geneseqp/AA1996.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cgdata/geneseq/geneseqp/AA1994.DAT:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cgdata/geneseq/geneseqp/AA1993.DAT:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cgdata/geneseq/geneseqp/AA1990.DAT:
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                                                R91438
W00859
                                                                                   R20811
R22550
                                                                                                                                                      W60555
R22549
                                                                                                                                    R91439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /geneseq/geneseqp/AA1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /geneseq/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /geneseq,
                                                                                                                                                                                                                                                    SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /geneseqp/AA1988.DAT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /geneseqp/AA1984.DAT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /geneseqp/AA1992.DAT:
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Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   390729
                                                                                                                                                                                                     Description
                                                                  Human FCRI (CDNA c
                                                                                                 Human macrophage-s
Human macrophage-s
                                                                                                                                                                    Oligopeptide from
                                    Human
                                                                                   Human macrophage-:
                                                                                                                                    Human FCRI (cDNA c
                                                    FCRI (CDNA C
Fc receptor
                 Fc receptor
                                                                                                                                                    macrophage-s
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	34	34	. 35	. 35	35	35	35	35	35	35	35	35	35	35	35	36	36	36	37	38	38	38	<u>အ</u>	38	38	38	38	38	53	55	56	61	61	61
•	55.7	55.7	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	. 59.0	59.0		60.7	62.3	. 62.3	62.3	62.3	62.3	62.3	62.3	٠		•	•	91.8	•	100.0	100.0
	124	38	1411	1193	365	293	293	249	249	201	201	201	170	129	129	838	-319	127	401	3080	. 803	793	793	793	793	, 787	787	. 787	377	261	410	399	374	374
	21	21	17	21	21	21	21	21	21	21	21	21	21	21	21	20	20	20	19	14	20	22	20	20	19	22	20	19	20	20	12	21	21	21
	G12546	B51764	W02258	Y90279	G14485	G21246	G14486	G21247	G14487	G35900	G21248	G10127	Y90275	G35901	G10128	W72975	Y41438	Y73820	W68417	R35081	Y43263	B28645	W81065	Y43262	W37911	B28646	W81064	W37912	W86195	.¥33183	R12428	B43683	δ	Y96134
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	Zea mays protein	Human secreted pro	Nucleolar/endosoma	Protein containing	Arabidopsis th			Arabidopsis thal	Arabidopsis thal	Arabidopsis th	Arabidopsis th	Arabidopsis th	PVBV NIa-pro p	Arabidopsis th			Fragment of human	Human prostate tum	Dengue virus type	ZYMV polyprotein	Human smoothened		Amino acid sequenc	w	Rattus norvegicus	Human Smoothened	Amino acid sequenc	₩.	Human Fc receptor	Human sFc-gammaRI		_		Human macrophage-
	in f	pro	oma	ling	thalia	thalia	thalia	ilia	lia	thalia	thalia	lia	prote	thalia	lia	17 gH	an	t um	/pe	٠	d p	≅	lenc	p b	Suc	ď.	lenc	noot	Ö,	RI	)RI	assoc	nit	e-s

## ALIGNMENTS

PXX W60555 De Kruif CA, Logtenberg T; 08-OCT-1996; 07-OCT-1997; 16-APR-1998. WO9815833-A1 Synthetic. cell transformation; mutation; anti; oncogene. Extracellular domain; CD64; identification; antibody; Oligopeptide from extracellular domain of CD64 18-AUG-1998 W60555 W60555 standard; peptide; 12 (UYUT-) RIJKSUNIV UTRECHT. lmmunohistochemical; (first entry) 96EP-0202791. 97WO-NL00557. immunofluorescent analysis; detection; B

WPI; 1998-240964/21.

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo; peptide derived from the target, useful for, e.g. identifying specific antibodies

Example 1;

Page

29;

40pp; English

Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a method for the identification of:a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the target protein and inable displaying the peptides.

epitopes on different molecules. The genes/olion selected peptides can be isolated and used for

The genes/oligonucleotides that encode

0;

recombinant production of for immunohistochemical or

antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent analysis, and also to detect cell transformation caused

by mutation in (anti)oncogenes.

that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two

surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments,

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RESULT
R22549
                                                                                                                                                                                                                                                                                                                                                                                                              망
                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                       WPI; 1992-056864/07
                                                         Seed B,
          New CD53 cell surface antigen
                              N-PSDB;
                                                                            (GEHO-)
                                                                                               13-JUL-1990;
                                                                                                               15-JUL-1990;
                                                                                                                                   23-JAN-1992
                                                                                                                                                     WO9201049-A
                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                       Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor
                                                                                                                                                                                                                  Human macrophage-specific FcRI receptor encoded by clone p98.
                                                                                                                                                                                                                                      21-MAY-1992 (first entry)
                                                                                                                                                                                                                                                            R22549;
                                                                                                                                                                                                                                                                             R22549 standard; Protein;
 immuno-therapy and diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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2
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| 1 ||qrpg|qlyfs | 12
                                                                                                                                                                                                                                                                                                                                           1 LLQRPGLQLYFS 12
                                                                           GEN HOSPITAL
                                                         Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                              12 AA;
                                                                                              90US-0553759
                                                                                                               90WO-US04986
                                                         Amiot M
                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                            344 AA.
                                                                                                                                                                                                                                                                                                                                                               0
of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                      Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                      9.5e-05;
                                                                                                                                                                                                                                                                                                                                                                                 DB 19;
                                                                                                                                                                                                                                                                                                                                                             0;
it - for
neoplasms,
                                                                                                                                                                                                                                                                                                                                                                              Length 12;
                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                            Gaps
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This amino acid sequence was predicted from the CDNA sequence of CDNA clone p98/X2. It differs from the sequence predicted from clone p185 (see Q211/78) at position 25, p135 encodes a Ser reside while clone p98/X2 predicts a Thr residue. At position 58, p90

Ser residue 58, p90

The amino acid sequence (CDNA clone, p98 (714718), obtd. from a cDNA library using an cDNA clone, p98 (714718), obtd. FCRI is a high affinity receptor immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (714717) and p90 (714719), coded for a variants of the FCRI sequence; the C-terminal sequence coded for a variants of the FCRI sequence with those of the p135

Example 10; Column 55-56; 79pp; English

sequence (R91439) of human

Cloning of cDNA encoding cell surface antigen – useful for isolation of diagnostic and therapeutic proteins

Example 10;

Page 94a;

160pp;

English.

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RESULT
R91439
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                                                                                                                                                                                                                                                                                                                                                       8888888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 100.0%; Score 61; DB 13; Best Local Similarity 100.0%; Pred. No. 0.0027; Matches 12; Conservative 0; Mismatches 0
                                                                                                                                                                                                                     01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                           N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                            Aruffo A,
                                                                                                                                                                                                                                                                                       25-FEB-1988;
                                                                                                                                                                                                                                                                                                            09-APR-1996
                                                                                                                                                                                                                                                                                                                                  US5506126-A
                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                            Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-0CT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R91439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R91439 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                 (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 LLQRPGLQLYFS
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                                                                                                                                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                             T14718
                                                                                                                                                                             Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                     92US-0983647.
88US-0160416.
89US-0379076.
89US-0553759.
93US-0139273.
                                                                                                                                                                                                                                                                                      88US-0160416
                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers 58
                                                                                                                                                                                                                                                                                                                                                                 /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12
                                                                                                                                                                                                                                                                                                                                                                                                                                vector;
                                                                                                                                                                                                                                                                                                                                                                                                                               cloning; immunoselection;
ector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                    "amino acid 58 is Leu
translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                344 AA
                                                                                                                                                                                                                                                                                                                                                                in
                                                                                                                                                                                                                                                                                                                                                                clone
                                                                                                                                                                                                                                                                                                                                                                                                                                        immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                               p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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0;

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RESULT
Y96183
ID Y9
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Best Local S
Matches 12
                                                                                                                                                                     /note-
Misc-difference 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and p90 products (see also R91438 and W00859). A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                      /note= "encoded
Misc-difference 126
                                                                                                                                                                                                                 /note-
Misc-difference 116
                                                                                                                                                                                                                                                                                      /note=
Misc-difference 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
Misc-difference
                                                          /note=
Misc-difference 136
                                                                           /note-
Misc-difference 134
                                                                                                /note= "encoded
Misc-difference 129
                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                   /note=
Misc-difference 82
                                                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                      Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human macrophage-specific FcRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-DEC-2000 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y96183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Y96183 standard; Protein; 344 AA
                    Misc-difference
                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Macrophage; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         y match 100.0%;
Local Similarity 100.0%;
hes 12; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 LLORPGLOLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 344 AA;
                                                                                                                                                                                                                                                                                                                                   /note-
55
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                                                                                                                                                        /note=
123
                                                                                                                                                                                                  /note= "encoded
117
                      /note=
140
                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                      /note=
                                                                                                                                                                                                                                                                               'note=
                                                                                                                                                                                                                                                                                                                         note=
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                                                                                                                                                                                                                                                                                                                                                                                            'note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cell surface antigen;
                                                                                                                                                                                                                                                         "encoded by CAG"
                                                                                                                                                                                                                                                                                                                         "encoded by CAC"
                                                                                                                                                                                                                                                                                                                                                                      "encoded by CGG"
         "encoded by AAT"
                               "encoded
                                                     "encoded
                                                                                                                                                                   "encoded
                                                                                                                                                                                        "encoded
                                                                                                                                                                                                                                    "encoded
                                                                                                                                                                                                                                                                             "encoded by CCC"
                                                                                                                                                                                                                                                                                                   "encoded by
                                                                                                                                                                                                                                                                                                                                               encoded by
                                                                                                                                                                                                                                                                                                                                                                                           "encoded by ACC"
                                                                                                                                                                                                                                                                                                                                                                                                                 "encoded by GCG"
                                                                           "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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Pred. No. 0.0027;
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                                                                                                                                                                by TTC"
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                                                                                                                                                                                                                                                                                                                                               ACC "
                                                                           AAT"
                                                                                                                                              ACC"
                                                                                                                                                                                         GCA"
                                                                                                                                                                                                                                                                                                     TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Ş

1 LLQRPGLQLYFS 12

Query Match Best Local S Matches 12

Similarity

100.0%;

Score 61; Pred. No.

DB 21; 0.0027;

Length 344; Indels

0

Gaps

0;

0;

Mismatches

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13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note-
Misc-difference 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                          The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US6111093-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
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Sequence
                                                               protein of
                                                                                                                                                                                       treatment of immune mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,
                                                                                                                                                                                                                                                                          CSA nucleic acids isolated by the method of the invention, and
                                                                                                                                                                                                                                                                                                                                       based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                           the invention designed to isolate CSA nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Column 53-55; 75pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                  surfaces other than macrophages,
                                                                                                      ability to interfere with such bonding,
                                                                                                                          immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The
                                                                                                                                                                                                                                                    proteins they encode, are useful
                                                                                                                                                                                                                                                                                                                   physica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-OCT-1998;
                                                                                                                                                                                                                              immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                              panning on) an antibody-coated substrate such as a culture
                                                                                                                                                                                                                                                                                                                 selection of cells expressing the antigen by adhesion
    344
                                                               FcRI and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92US-0983647.
88US-0160416.
89US-0379076.
89US-0498809.
90US-04553759.
  ΑĄ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seed B;
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216
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338
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333
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306
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305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-
220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note=
                                                             a receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "encoded by GTG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "encoded by CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded_by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded by CGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded by
                                                               ligand will
                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
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                                                                                                                                                                                                                                                    for immunodiagnostic and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT"
                                                             ces of macrophages. The or to cause it to occur on ful in therapy. A fusion ill be helpful to increase
                                                                                                                                                                                                                                                                                                                                                         The method
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217 llgrpglqlyfs 228

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                                                                                                                            RESULT
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                                                                                                                                                                                                                                                                 Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 12; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                    cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion prof PCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This amino acid sequence was predicted from the cDNA sequence of cDNA clone pl35. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25, pl35 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, pl35 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the
                                            R22550;
                                                                                R22550 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane spanning domain, and a short, highly charged
  21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-MAY-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R20811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R20811 standard; Protein; 374 AA
                                                                                                                                                                               1 LLQRPGLQLYFS 12
||||||||||
|217 llqrpglqlyfs 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W09201049-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                          σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                    .374 AA
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90WO-US04986
                                                                                  Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Amiot M
                                                                                  374
                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                 Score 61; DB 13;
Pred. No. 0.003;
); Mismatches 0;
                                                                                  A
                                                                                                                                                                                                                                                                                                                                                                                                                                                Ig-related domains. A fusion protein
                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                       Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     etc.
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RESULT
R91438
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and clone p90 predicts a Thr residue. At position 58, p90 predicts, val while clones p135 and p98/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-relate domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                     Key
Misc-difference
                                                                                                                                                                            Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                              Human FcRI (cDNA clone pl35 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue
                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                    30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                    R91438;
                                                                                                                                                                                                                                                                                                                                                                                                  R91438 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rapid immunoselection cloning technique; cell surface antigen, immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human macrophage-specific FcRI receptor encoded by clone p90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    217 llqrpglqlyfs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 LLQRPGLQLYFS
                                                                                                                            sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ΑĄ;
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                                                     Location/Qualifiers 25
/note= "amino acid 25 is Thr in clone p90 and p98
translated products"
                                                                                                                                                                                                                                                                                                                                                                                               Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                    374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 61; DB 13; Length 374; Pred. No. 0.003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                 immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Misc-difference 58

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/note=

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Best Local Similarity 100 Matches 12; Conservative
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Best Local
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The amino acid sequence (R91438) of human FcRI was detd. from a cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity receptor for the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contract of the contrac
                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W00859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cloning of cDNA encoding cell surface antigen - useful of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                 09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cell surface antigen; cloning; immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W00859 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FcRI sequence, and a third clone, p98 (T14718), coded for an FcRI (R91439) having a different C-terminal sequence. A fusion protein of FcRI and a receptor ligand will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-APR-1996
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                                                                                                                                                        25-FEB-1988
                                                                                                                                                                                                                                                                                                         US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 55-56; 79pp; English:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        æ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    llqrpglqlyfs 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                        88US-0160416
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900S-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-0983647.
88US-0160416.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 61;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 374;
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RESULT
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Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The amino acid sequence (W00859) of human FCRI was detd. from a cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity recepts for the FC portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                        Misc-difference
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                                                                                                                                                                                                                                                                                                                               cloning.
                                                                                                                                                                                                                                                                                                                                                                     Human Fc receptor I.
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                                                                                                                                                                                                                                                                                                                                                                                                                        W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                W80448 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; T14719
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18-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GEHO ) GEN
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||||||||||
| 217 llqrpglqlyfs 228
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93US-0139273.
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60
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159
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                                                                                                                                                                                                                                                                                                                                         cell surface antigen; lymphocyte; human;
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            "encoded by
                                                                                                              "encoded by AAT"
                                      "encoded
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Pred. No.
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            GGC
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                                                                CAA
                                                                                                                                           CCC"
                                                                                                                                                                    CTG"
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Misc-difference

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W97833
                                                       RESULT 10
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Best Local
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
  W97833;
                                                                                                                                                                                                                                                                                 produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                  expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell ratace has been used to clone genes (see V6342-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
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                            W97833 standard;
                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                         immunotherapeutic applications, treatment of immune-mediated in
                                                                                                                                                                                                                                                                                                                                          genes can
                                                                                                                                                                                                                                                                                                                                                                                                                                                 from a single patient undergoing extracorporeal induction therapy. Clones p90 (see X07372) and were also obtained. A novel method for cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This is the amino acid sequence of human Fc receptor I (FCRI), deduced from cDNA clone pl35 (see V63456) isolated using a rapi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-NOV-1998
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                                                                                                                                                                                                                                                                                                                          genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunoselection cloning method from a cDNA library expressed
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                                                                                              217
                                                                                           llqrpglqlyfs 228
                                                                                                                        LLORPGLOLYFS 12
                                                                                                                                                  l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The cDNA library was constructed from polyA RNA of cells
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                                                                                                                                                 100.0%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                           immune-mediated infections, diseases,
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89US-0379076.
90US-0553759.
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171
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176
                          Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         75pp;
                             374 AA
                                                                                                                                                  0;
                                                                                                                                                            Score 61; DB 19;
Pred. No. 0.003;
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                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNAS
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                                                                                                New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
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Misc-difference 159
induction therapy. were also obtained. expression librarie
                                                     This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                             Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                           03-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Fc
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              from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mammalian
                                                                                                                        Example 10; Column 53-54; 75pp;
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DB; x07372.
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    libraries
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23
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176
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141
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    is based on
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                        Misc-difference
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Misc-difference 171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W97834 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
US5830731-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human Fc receptor I
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treatment of immune-mediated infections,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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ilarity 100.0%;
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• 159
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256
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Pred. No. 0.003;
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the antigen by is useful for
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13-JUL-1990;
21-MAY-1997;
                                                                          has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                             expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                             This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p98/x2 (see x073/73) isolated using a rapid immunoselection cloning method from a cDNA library expressed in Cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
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25-FEB-1988;
                                        treatment of immune-mediated infections, animals, including humans.
                                                                                                                                                                                                                                                                                                                                                   were also obtained.
                                                                                                                                                                                                                                                                                                                                                                     from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p90 (see X07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; Column 53-54; 75pp; English.
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DB; X07373.
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  374 AA;
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90US-0553759
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88US-0160416
                                                                                                                                                                                                                                                                                                                                                   Clones p135 (see V63456) and p90 (see X07372) A novel method for cloning cDNAs from mammal
                                                              and disorders of
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В Ş Query Match
Best Local Similarity
Matches 12; Conserv 1 LLQRPGLQLYFS 12 100.0%; ilarity 100.0%; Conservative 0, Score 61; DB 19 Pred. No. 0.003; Mismatches

DB 19;

Length 374; Indels

0;

Gaps

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RESULT 12 Y96134 19-DEC-2000 Y96134; Human macrophage-specific FcRI. Y96134 standard; Protein; 374 AA (first entry)

XXXXX

amyloidosis; Macrophage; FcRI; cell surface antigen; human; immunoselection; immune panning; disorder; immunodiagnosis; diagnosis; infection; asthma; ma; immunotherapy; gene th
ma; immune-complex disease;
parasitic disease; Fc rece therapy;

2000-586382/55 DB; A50592.

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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
Stamenkovic I,
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                 (GEHO ) GEN HOSPITAL CORP
                                                                                    28-OCT-1998;
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89US-0379076.
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90US-0553759.
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                                                                                   9808-0181612.
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RESULT
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Best Local
            WPI; 2000-367968/32
N-PSDB; A27466.
                                                           (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                           Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IGG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                             03-DEC-1998;
                                                                                                                              03-DEC-1998;
                                                                                                                                                              07-JUN-2000.
                                                                                                                                                                                             EP1006183-A1
                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                               systemic lupus
                                                                                                                                                                                                                                                                                                                                                               Human high affinity Fc receptor,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     surfaces other than macrophages, protein of FcRI and a receptor 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ability to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   proteins they encode, are useful immunotherapeutic applications, i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is that of a human macrophage specific PcRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated nucleic acid molecule encoding the CD19 cell surface a useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, p.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  treatment of immune-mediated infections, diseases, and diso animals, including humans. These disorders include asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         physical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     based
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    physical selection of cells expressing the antigen by adhesion to panning on) an antibody-coated substrate such as a culture dish.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     potencies of antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  upon transient expression of a CSA
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12; Conserv
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ther than macrophages, is useful
FCRI and a receptor ligand will
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Column 53-55; 75pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                               erythematosus; tumour.
                                                                                             98EP-0122969
                                                                                                                              98EP-0122969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 61; DB 21
Pred. No. 0.003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l for immunodiagnostic and including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                               FcgammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    diseases, and disorders in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in therapy.
be helpful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 374;
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  DRX PRX FXX
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                may be modified to produce recombinant versions. The recombinant receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cells expressing Fc receptors. FcgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence is the human high affinity Fc receptor, FcgammaRI FcgammaRI is also known as CD64. Fc receptors play an important
                                                                                                                                                                                                                                                                                                                                                                                            immune disorder; haematopoietic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        B43683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B43683 standard; Protein; 399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel Fc receptor lacking transmembrane domains, glycosylation, useful for diagnosing and treatin cancer -
WPI; 2000-587533/55
                                                                                                                                                                          08-MAR-2000; 2000WO-US05882
                                                                                                                                                                                                                      21-SEP-2000
                                                                                                                                                                                                                                                                 WO200055350-A1
                                                                                                                                                                                                                                                                                                                                                     neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                     dermatological; neuroprotective; thrombolytic; coagulant; nootropic;
vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human cancer associated protein sequence SEQ ID NO:1128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                             Rosen CA,
                                                                                    (HUMA-) HUMAN GENOME
                                                                                                                               12-MAR-1999;
                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; cancer associated gene; cancer antigen; detection;
                                                                                                                                                                                                                                                                                                                                                                            haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 LLQRPGLQLYFS 12
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|217 llqrpglqlyfs 228
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l Similarity 100.0%;
l2; Conservative (
                                             Ruben SM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic;
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                                                                                                                                                                                                                                                                                                                                                   thrombolytic;
disease; drug
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                                                                                                                               99US-0124270.
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                                                                                      SCI INC
                                                                                                                                                                                                                                                                                                                                                     drug screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              proliferative; vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60pp; English
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                            disorder; infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunomodulator;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer;
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                                                                                                                                              PXPX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and 194240 represent sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neuroprotective; cardiant; thrombolytic; coaquiant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating med conditions and diagnosing pathological conditions. Polynucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C77607 to C78448 encode the human cancer associated proteins B43398 to B44239. The proteins can have activities based on and cells the genes are expressed in Example of activities
                                                                                                                                                                                                                                                                                                                                                                                                                       Hybrid Fc(gamma)RII/I receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R12428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R12428 standard; Protein; 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytostatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g.\ 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N-PSDB; C77892
WPI; 1991-164135/22
                                    Hogarth PM,
                                                                                                                                                                                  16-MAY-1991
                                                                                                                                                                                                                                                                                             Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                      (UYME-) UNIV MELBOURNE
                                                                                                          25-OCT-1989;
                                                                                                                                              25-OCT-1990;
                                                                                                                                                                                                                       WO9106570-A
                                                                                                                                                                                                                                                                                                                               Region
                                                                                                                                                                                                                                                                                                                                                                                   Immunoglobulin-binding molecule; Fc(gamma)RI; Fc(gamma)RII
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12; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proliferative; vulnerary; immunomodulator; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                    Hulett MD,
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                                                                                                          89AU-0007045
                                                                                                                                                90WO-AU00513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    encode the human cancer associated proteins given in . The proteins can have activities based on the tissues
                                                                                                                                                                                                                                                                                               204..410
                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                            /label=
                                                                                                                                                                                                                                                                                                            /label= Fc(gamma)RII domains dl and
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                                                                                                                                                                                                                                                                          D3,
                                      McKenzie IFC,
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                                                                                                                                                                                                                                                                        transmembrane and
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DR N-PSDB; Q12079.

XX

PT New immunoglobulin binding hybrid Fc receptor molecules - used to PT control auto-immune diseases and allergic reactions and to regulate. PT antibody prodn.

XX Claim 10; fig 3B; 80pp; English.

PS Claim 10; fig 3B; 80pp; English.

CC This chimeric sequence is one example of a hybrid Fc receptor CC of the invention. The hybrid Fc receptor retains the Ig binding CC functions of the constituent receptor types. It can be used for CC regulating antibody production in vivo and in plasmaphoresis to CC remove immune complexes or pathological antibodies. Rabbit IgG CC bound to cells transfected with the chimeric Fc(gamma) receptor CC See also Q12078.

XX Squence 410 AA;

Query Match 91.8%; Score 56; DB 12; Length 410; Best Local Similarity 91.7%; Pred. No. 0.026; Matches 11; Conservative 1; Mismatches 0; Indels 0; Gaps 0; Ulioptical Stationary of the completed: June 4, 2001, 12:13:28

Qy 1 LLORPGLOLYFS 12

Db 232 llqrpg1qlhfs 243

Search completed: June 4, 2001, 12:13:28
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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Perfect score:
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Gapop 10.0 , Gapext 0.5
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
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PCT-US95-13703-21

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US-09-293-395-13

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US-09-293-395-13

US-09-293-395-13

US-07-857-224B-48
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PCT-US95-08950-12
US-07-952-853-22
US-08-914-848-22
US-08-840-316-3
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•	US-08-936-135-4	US-09-102-248-5	US-08-801-263A-5	US-09-102-248-9	US-08-801-263A-9	US-09-102-248-2	US-08-801-263A-2	US-09-090-793-9	US-08-752-929-15	US-08-375-709-15	PCT-US94-05277-2	US-09-074-658-19	US-08-867-941-19	US-08-613-009A-7	US-08-613-009A-11	US-08-613-009A-8	US-08-630-916A-48	US-09-357-746-4
	Sequence 4, Appli	Sequence 5, Appli	Sequence 5, Appli	Sequence 9, Appli	Sequence 9, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 9, Appli	Sequence 15, Appl	Sequence 15, Appl		Sequence 19, Appl	Sequence 19, Appl	Sequence 7, Appli	Sequence 11, Appl	Sequence 8, Appli	Sequence 48, Appl	Sequence 4, Appli

#### ALIGNMENTS

US-08-332-562A-94

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TOPOLOGY:
US-08-332-562A-94
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                                                                                                      TELEX: 904136
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                             FILING DAIL:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/896,457
APPLICATION NUMBER: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                               SEQUENCE CHARACTERISTICS:
-LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                            FILING DATE: 27-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54270/119/GRHA
                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Foley & Lardner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HULETT, Mark D.
TITLE OF INVENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
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20007-5109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HIBBS, Margaret L. SCOTT, Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HOGARTH,
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HOGARTH, Mark P.
                             single-
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Best Local Similarity

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JS-08-953-823A-4
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US-08-720-484A-4
                                                                                                                                                                                  Sequence 4, Application US/08953823A Patent No. 6136958
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                                                                                                                                                                    ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NFORMATION FOR SEQ ID NO:
                                                                                APPLICANT: DeSauvage; Frederic
APPLICANT: Rosenthal, Arnon
APPLICANT: Stone, Donna
TITLE OF INVENTION: Vertebrate
NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 787 amino acids
                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                  579 LLQNPGQELSFS 590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WinPatin (Gen
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb |
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT: Rosenthal, Arnon
APPLICANT: Stone, Donna
APPLICANT: Stone, Donna
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 650/225-1489
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                                                                                                                                                                                                                                                                                                                                                     Local Similarity 66.
nes 8; Conservative
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                                 STREET:
                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Syoboda, Craig G.
REGISTRATION NUMBER: 39,044
                                                                                                                                                                                                                                                                                                      1 LLQRPGLQLYFS 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BER OF SEQUENCES:
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South San Francisco
California
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                                 E: Genentech, Inc
1 DNA Way
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1 DNA Way
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DN: 530
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Pred. No. 41;
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1; Mismatches 0;
                                                                                                Smoothened Proteins
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US-08-720-484A-2
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Best Local Similarity 66.7%;
Matches 8; Conservative
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Patent No. 5990281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 787 amino acids
TYPE: Amino Acid
                                                                                                                                          SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/720,484A
FILING DATE: 30-Sep-1996
                   TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1489
                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Svoboda, Craig G.
REGISTRATION NUMBER: 39,044
                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inc
                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Rosenthal, Arnon APPLICANT: Stone, Donna
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/00
FILING DATE: 30 -Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: Syoboda, Craig G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA
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MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PC_DOS/MS_DOS
                                                                                                                                                                                                                                                                                                                        STREET: 1 DNA TO CITY: South San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  579 LLQNPGQELSFS 590
                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                  CLASSIFICATION:
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Syoboda, Craig G.
Syoboda, Craig G.
Syoboda, Craig G.
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650/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DeSauvage, Frederic
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                                                                                                                                                                                                                                                                3.5 inch, 1.44 Mb floppy disk
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30-Sep-1997
V: /
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TYPE: P

1: 793 amino acids Amino Acid

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Query Match
Best Local Similarity
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Sequence 4, Application US/08216260 Patent No. 5837261
                                                                                                                                                Query Match
Best Local Similarity Matches 8; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/08953823A Patent No. 6136958
                                                                                                                                                                                                                                                                                                   NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/027070
FILING DATE: 30-Sep-1996
                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 39,044
REFERENCE/DOCKET NUMBER: P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1489
                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                        583 LLQNPGQELSFS 594
                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 30-Sep-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/953,823A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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                                                                                                                    1 LLORPGLOLYFS 12
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                                                                                                                                                                                                                                                     4: 793 amino acids
Amino Acid
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                                                                                                                                                                                                                                                                                                                                                                                      Svoboda, Craig G.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DeSauvage, Frederic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WinPatin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM PC compatible SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-Sep-1997
                                                                                                                                                               66.7%;
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                                                                                                                                                               Score 38; DB 4;
Pred. No. 41;
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Pred. No.
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5304637-7
                                                                                                                                                 Query Match

Best Local Similarity

Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PATENTIN RELE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Minson, Anthony C. PITLE OF INVENTION: VIRAL VACCINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 19-MAR-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION DATA:
                                                                             227 LLRSPGRYVYFS 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/168,643 FILING DATE: 16-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/030,073 FILING DATE: 19-MAR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: FILING DATE: 06-DE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: United States ZIP: 94111-4187
                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                             1 LLQRPGLQLYFS 12
                                                                                                                                                                                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           California
                                                                                                                                                                                                                                                                                                                                                                        910 277299
                                                                                                                                                                                                                                                                                                               838 amino acids
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                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inglis, Stephen C
                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Flehr, Hohbach, Test, Albritton & Herbert Attn: Walter H. Dreger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-DEC-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-MAR-1993
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                                                                                                                                                   Mismatches
                                                                                                                                                                  Score 36; DB 2;
Pred. No. 1e+02;
                                                                                                                                                                                       DB 2;
                                                                                                                                                                                   Length 838;
                                                                                                                                                   Indels
                                                                                                                                                   0;
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APPLICANT: DORSSERS, LAMBERTUS C.J.; WAGEMAKER, GERARD; VOS, VONNE J.; VAN LEEN, ROBERT W.; PERSOON, MARIA L.N.
TITLE OF INVENTION: EXPRESSION AND PURIFICATION OF HUMAN

0

INTERLEUKIN-3 AND MUTEINS THEREOF
NUMBER OF SEQUENCES: 30
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/ FILING DATE: 13-MAR-1990

US/07/494,182

NUMBER: 249,184

Patent No. 5304637

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Best Local Similarity
thes 7; Conserv
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5304637-7
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5304637-1
;Patent No. 5304637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: DORSSERS, LAMBERTUS C:J.; WAGEMAKER, GERARD; VOS VONNE J.; VAN LEEN, ROBERT W.; PERSOON, MARIA L.N.
TITLE OF INVENTION: EXPRESSION AND PURIFICATION OF HUMAN INTERLEUKIN-3 AND MUTEINS THEREOF NUMBER OF SEQUENCES: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; FILING DATE: SEQ ID NO:7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12, Application US/08284393B
APPLICATION NUMBER: US/08/284,393
ETLING DATE: 01-AUG-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX0389
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       tent No. 569623
                                                                                                                                COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 249,184
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Zurawski, Sandra M.
APPLICANT: Zurawski, Gerard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                   ITLE OF INVENTION: MUTEINS OF MAMMALIAN CYTOKINES
                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                     CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 16-AUG-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 150
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                                                                                                                                                                                                                                     94304-1104
                                                                                                                                                                                                                                                                    California
                                                                                                                                                                                                                                                                                                 901 California Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DORSSERS, LAMBERTUS C.J.; WAGEMAKER, GERARD; VOS,
                                                                                                                                                                                                                                                      USA
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87.5%;
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87.5%;
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Pred. No.
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22;
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Best Local Similarity
7; Conserva
                                                                               PCT-US95-08950-12
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PCT-US95-08950-12
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                Query Match
Best Local (
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                                                                                                                                                                                      NEORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: 415-852-9196
TELEFAX: 415-496-1200
INFORMATION FOR SEQ ID NO:
                                                                                             MOLECULE TYPE:
                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 152 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Zurawski, APPLICANT: Zurawski,
                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 415-852-9196
                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Ching, Edwin P.
REGISTRATION NUMBER: 34,
                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
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APPLICATION NUMBER: U
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                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                          TOPOLOGY:
                                                                                                                         STRANDEDNESS:
                                                                                                                                           TYPE:
                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                 FILING DATE: 01-AUG-199
                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Palo Alto
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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    LLQRPGLQ 8

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: . DNAX Research Institute
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    Similarity 7; Conser
                                                                                                                                         amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               901 California Avenue
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                                                                                                                                                                                                                                                                                                                                                                                                                       PatentIn Release #1.0, Version #1.25
                                                                                                            linear
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87.5%;
              54.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MUTEINS OF MAMMALIAN CYTOKINES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sandra M.
                                                                                                                                                                                                                                                                                                                              US 08/284,393
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                                                                                                                                                                                                                                                                                                                                                                                          PCT/US95/08950
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Score 33; DB 5;
Pred. No. 57;
0; Mismatches
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Pred. No. 57;
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                              Length 152;
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Matches

Conservative

Indels

Gaps

0,:

12 LLVRPGLQ 19

1 LLQRPGLQ 8

RESULT 11 US-07-952-853-22

Application US/07952853

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; MOLECULE TYPE:
US-07-952-853-22
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Best Local Similarity
                                                                                                                 Sequence 22,
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                                                                                                  tent No.
                                                                                                                                                                                                                                                                                                                                                                                                        NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 628 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
                                                           APPLICANT:
                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US FILING DATE: 19921125 CLASSIFICATION: 435
                                                                                                                                                                                       62 MLQNPGLQ 69
                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Murashige, Kate
REGISTRATION NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Palo Alto
STATE: California
ZIP: 94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                      1 LLQRPGLQ 8
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                                                                                    INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                               amino acid
                                                                                                                 Application US/08914848
                                                                                                                                                                                                                                                                                                                                                                             628 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                      415-494-0792
                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            De Graaff, Leendert H
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                                                      Bakhuis,
                                                                       Van Heuvel, Margaretha
                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                                  protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
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, Michel J. A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jacob
                           Abraham
                                                                                                                                                                                                                                                             54.18;
75.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Janna G.
                                                        Janna G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peter M.
             Leendert H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US/07/952,853
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Pred. No. 2
                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                          DB 2;
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                                                                                                                                                                                                                                                                         Length 628
                                                                                                                                                                                                                                             Indels
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US-08-914-848-22
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Best Local Similarity
                                                                                                                                                                                                                                                                                            Sequence 3;
Patent No. (
                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 628 amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 24
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-813-5600
                                  COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                               TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 25-NOV-1992
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US 0
FILING DATE: 25-NOV-1992
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                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Call
STP: 94304-1018
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS MOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                   CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                       62 MLQNPGLQ 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX:
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                                                                                  COUNTRY:
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                                                                                                                                               ADDRESSEE:
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                                                                     10154
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                                                                                                                                                                                                                                                                                                         Application US/08840316
                                                                                                                                    345 PARK AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               628 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 755 Page Mill Road
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                     USA
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                                                                                                                                                   MORGAN & FINNEGAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM PC compatible SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MORRISON
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                                                                                                                                                                             Use In Diagnostic Methods and Vaccines
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75.0%;
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      PC-DOS/MS-DOS
                                                                                                                                                                                                                                                            Suzanne U., Purcell, Robert H.;
                                                                                                                                                                                                                A Pakistani Strain Of Hepatitis E And Their
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peter M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US 07/952,853
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 628
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CURRENT APPLICATION DATA:

APPLICATION NUMBER:

US/08/840,316

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Query Match
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Best Local Similarity bu...
Conservative
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APPLICATION NUMBER: US 08/279,823
FILING DATE: 25-UL-1994
PRIOR APPLICATION DATA:
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TELEPHONE: (212) 758-4800
TELEFAX: (212) 751-6849
NFORMATION FOR SEQ ID NO: 3:
                                                            PRIOR APPLICATION DATA:
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                                                                                                                                                                                                            URRENT APPLICATION DATA:
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                                                                             APPLICATION NUMBER: US 0 FILING DATE: 05-APR-1991
                                                                                                                                                                            APPLICATION NUMBER: US/08/478,507 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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83 RPGLDLVFA 91
                                APPLICATION NUMBER: US 07/505,888 FILING DATE: 05 APR-1990
                                                                                                                                                                                                                               SOFTWARE:
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                                                                                                                                                                                                                                                                                                              ZIP: '94306
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                                                                                                                                                                                                                                                                                                                                                         CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                          STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Richard W. Bork REGISTRATION NUMBER: 36 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 RPGLQLYFS 12
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                                                                                                                                                                                                                                                                                                                                              CA
                                                                                                                                                                                                                                                                                                                                                                        3: Dehlinger & Associates
350 Cambridge Avenue, Suite 250
                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                           PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tam, Albert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krawczynski, Krzysztof Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Yarbough, Patrice O
Bradley, Daniel W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Reyes, Gregory R
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                                                                                                                                                                                                                                                                           Floppy disk
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                                                                                           US 07/681,078
US 07/420,921
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36,459
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            equence 3, Application No. 6207416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NFORMATION FOR SEQ ID NO:
                                      APPLICATION NUMBER: US08/316,765
FILING DATE: 03-OCT-1994
PRIOR APPLICATION DATA:
                                                                                       FILING DATE: 03-OCT-PRIOR APPLICATION DATA:
                                                                                                                                  FILING DATE: 28-MAY-1
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ENERAL INFORMATION:
                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
                                                                                                                                                                                    CURRENT APPLICATION DATA:
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TTORNEY/AGENT INFORMATION:
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TOPOLOGY: linear
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ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
           APPLICATION NUMBER: 07/9
FILING DATE: 18-SEP-1992
                                                                                                     APPLICATION NUMBER: PCT/US95/13102 FILING DATE: 03-OCT-1995
                                                                                                                                                    APPLICATION NUMBER: US/08/809,523 FILING DATE: 28-MAY-1997
                                                                                                                                                                                                   SOFTWARE:
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                                                                                                                                                                                                                                                                                10154
                                                                                                                                                                                                                                                                                                         NEW YORK
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                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                              IBM PC COMPATIBLE SYSTEM: PC-DOS/MS-DOS WORDPERFECT 5.1
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                             07/947,263
                                                                                                                                                                                                                                                                                                                                                                                                                   Pakistani
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Pred. No.
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REFERENCE/DOCKET NUMBER: 2026-4032US4

TELECOMMUNICATION INFORMATION:
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A;Accession: I48471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fc gamma (IgG) receptor high affinity - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
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A; Residues: 1-336 < RES>
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Science 260, 695-698,
                                                                                                                                                                                                                                                                                                                                 RESULT
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#### ALIGNMENTS

Todd, J.A.; Rodrigues, N.R.;

Ghosh, S.;

Hogarth, P.M.; Wicker,

R;Allen, J.M.; Seed, B. 11824, 1988
Nucleic Acids Res. 16, 11824, 1988
A;Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (A;Reference number: S03018; MUID:89098339 Query Match Best Local Similarity Watches 12; Conserv. N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000-C;Accession: A41357; S03019 C;Keywords: immunoglobulin receptor F;128-180/Domain: immunoglobulin homology <IMM> A;Cross-references: EMBL:X70980; NID:g311748; PIDN:CAA50311.1; PID:g311749 C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: immunoglobulin receptor C;Keywords: A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Molecule type: mRNA A; Residues: 1-344 <ALL1> A; Reference number: A; Accession: A41357 A; Title: Isolation and expression A; Reference number: A41357; MUID: R; Allen, J.M.; Seed, I Science 243, 378-381, A; Accession: Fc gamma (IgG) receptor I (high affinity) 231 LLQRPGLQLYFS 242 1 LLQRPGLQLYFS 12 references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 S03019 1135, Seed, B. 281, 1989 Conservative 100.0%; MUID:89100284 0; of functional high-affinity Fc Score 61; Pred. No. Mismatches form DB 2; ۵ .00096; Length 336; Indels receptor complementa 0; Gaps .0

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A; Note: the authors translated the codon ACO C; Superfamily: Fc gamma receptor I; immunoglobu C; Keywords: alternative splicing; immunoglobu
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Best Local Similarity 100.
The Conservative
               ;Superfamily: Fc gamma receptor I; immunoglobulin homology ;Keywords: alternative splicing; glycoprotein; immunoglobulin ;1-15/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                               ;Molecule type: mRNA;Residues: 1-24,'S',26-337,'T',339-374 <AL2>;Residues: 1-24,'S',26-337,'T',339-374 <AL2>;Cross-references: EMBL:X14356; NID:g31331; PIDN:CAA32537.1; PID:g31332;Perez, C.; Wietzerbin, J.; Benech, P.D.
ol. Cell. Biol. 13, 2182-2192, 1993
ol. Cell. Biol. 13, 2182-2192, 1993
;Title: Two cis-DNA elements involved in myeloid-cell-specific expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change; Accession: A39878; I70304; B41357; S03018; I57525; Accession: Anderson, C.L.; Chiu, I.M.; Van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M.; Biol. Chem. 266, 13449-13455, 1991
                                                                                                                                                                                            Residues: 1-8 <RE2>
                                                                                                                                                                                                                                       Status: translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Title: Nucleotide sequence of three cDNAs Reference number: S03018; MUID:89098339
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                                                                                                                                                                     Cross-references: GB:S57204; NID:g298692;
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cleic Acids Res. 16, 11824,
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coss-references: GB:X14356; GB:M21091; NID:g31331;
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MUID:93055454
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Pred. No. 0.00098;
); Mismatches 0;
predicted <EXT>
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                                    receptor;
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N;Alternate names: nign arrange 21-Jan-2000 C;Species: Mus musculus (house mouse) C;Species: Nis musculus (house mouse) C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jun-1994 #text_change 2
                                                                                                                                                                                                                                                                                          R;Krasney, P.A.; Carr, C.; Cavener, D.R.
Mol. Biol. Evol. 7, 155-177, 1990
A;Title: Evolution of the glucose dehydrogenase
A;Reference number: A39019; MUID:90205602
                                                                                                                                                                                                                                                                                                                                                                                                                                           glucose dehydrogenase (acceptor) (EC 1.1.99.10) - 1
C;Specias: Drosophila melanogaster
C;Date: 27-Nov-1991 #sequence_revision 09-May-1997
C;Accession: A39019; S06628
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F;293-313/Domain: transmembrane #status predicted <TMM>
F;293-315/Domain: transmembrane #status predicted <TMM>
F;59,78,152,159,163,195,240/Binding site: carbohydrate (Asn) (covalent) #status predicted (Asn)
                                                                          R;Whetten, R.; Organ, E.; K
Genetics 120, 475-484, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: immunoglobulin receptor; transmembrane protein F;127-179/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Structure and mapping of the gene encoding mouse A;Reference number: A46480; MUID:92166399 A;Accession: A46480
                               A; Title: Molecular structure and
                                                                                                         A;Cross-references: GB:M29298; NID:g157499; R;Whetten, R.; Organ, E.; Krasney, P.; Cox-
                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-612 <KRA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A39019
                                                                                                                                                                                                                                                               A; Accession: A39019
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Note: sequence extracted from R; Sears, D.W.; Osman, N.; Tate, J. Immunol: 144, 371-378, 1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fc gamma (IgG) receptor high affinity - mouse
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ilarity 100.0%;
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91.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI backbone (NCBIN:85205, NCBIN:85208, B.; McKenzie, I.F.C.; Hogarth, P.M.
transformation of the :89065357
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Pred. No. 0.00
1; Mismatches
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Pred. No.
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                                                                                                      7499; PIDN:AAA28571.1; Cox-Foster, D.; Caveno
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0.0097;
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                                   glucose dehydrogenase
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g syndromes.
F;1-32/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                Biochem. Biophys. Res. Commun. 235, 142-147, 1997

A;Title: Cloning of a mouse Smoothened cDNA and expression water and a service of a mouse smoothened cDNA and expression water.
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F;66-95/Region: beta-alpha-beta
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A;Cross-references:
                                                                     A; Experimental source: ADTC5
                                                                                  A; Molecule type: protein
A; Residues: 528-533;539-545;600-605 <AK2>
                                                                                                                                         A;Molecule type: mRNA
A;Residues: 1-793 <AK
                                                                                                                                                                        A; Accession: JC5539
                                                                                                                                                                                                                                                                                                                Smoothened protein precursor - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to the EMBL Data Library, July 1999
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                                                                                                                                                                                          A; Reference number: JC5539;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nypothetical protein PAB1688 -
                                                                                                                                                                                                                                                           ;Species: Mus musculus (house mouse);Date: 02-Sep-1997 #sequence_revision;Accession: JC5539; PC4476
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                                                                                                                        Accession: PC4476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gene: PAB1688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GB:AJ248286; GB:AL096836; NID:g5458366; PIDN:CAB49934.1; PID:g54584
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Residues: 1-393 <KAW>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Accession: A75079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   anonymous, Genoscope
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                 Comment: This protein is used in the conserved targets in Hedgehog signalling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Experimental source: strain
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77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 38;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred.
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Best Local Similarity
Matches 5; Conserv
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Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617
A;Accession: C64829
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Best Local Similarity
Thes 5; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Wu, M.; Yeong, C.Y.; Lee, S.C.; Wong, S.M. Nucletc Acids Res. 21, 1317, 1993
A;Title: Nuclectide sequence of the 3' half of A;Reference number: S30236; MUID:93219099
C;Superfamily: hypothetical protein b1328
C;Keywords: DNA binding; transcription re
F;19-49/Region: helix-turn-helix motif
                                                                                                                                                        A: Molecule type: DNA
A: Residues: 1-302 <BLAT>
                                                                                                                                                                                                                                                                                                                        R; Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; A.; Rose, D.J.; Mau, B.; Shao, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Note: the nucleotide sequence was submitted to the EMBL Data Library, C; Superfamily: tobacco etch virus genome polyprotein
                                                                                                               A;Cross-references: GB:AE000192; GB:U00096; NID:g1787125; PIDN:AAC73986.1; PID:g17871
A;Experimental source: strain K-12, substrain MG1655
                                                                                                                                                                                                                                                                                                                                                                      C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                              C;Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 29-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                      C; Species:
                                                                                                                                                                                                                                                                                                                                                                                                                                       transcription regulator yeaN - Escherichia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C64829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F;495-1010/Product: RNA-directed RNA polymerase #status predicted <POL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                F;62-494/Product: nuclear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   F;9-61/Product: VPg protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C; Keywords: coat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: mRNA
A; Residues: 1-1016 <WUM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C:Date: 19-Mar-1997 #sequence_revision 29-Aug-1997 #text_change 17-Nov-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Variety:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Species: zucchini yellow mosaic virus, ZYMV
                                                                                                                                                                                                         A; Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT
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                                                                   A; Gene:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cross-references: EMBL:X68509; NID:g288233;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Status: nucleic acid sequence not shown; translation not shown
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 179 IINKPGIQAYF 189
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8; Conserv
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protein; nuclear inclusion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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inclusion protein a #status predicted <NIA>
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                Perna, N.T.; Burland, V.; Riley,
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Length 302;

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G-protein signaling regulator - fission yeast (Schizosaccharomyces pombe) C;Species: Schizosaccharomyces pombe C;Date: 16-May-1996 *sequence_revision 13-Mar-1997 *text_change 21-Jan-2000 C;Accession: T38179; S62427 R;Lye, G.; Churcher, C.M.; Barrell, B.G.; Rajandream, M.A.; Walsh, S.V.
                                                                                 RESULT
S62427
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A;Residues: 1-307 <STO>
A;Cross-references: GB:AP001508; GB:BA000004; NID:g10172890;
A;Experimental source: strain C-125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypothetical protein F45Cl2.l4 - Caenorhabditis elegans
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A;Accession: H83705
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                                                                                                                                                                                                                                                                                                \;Introns: 27/2; 337/3
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Johnson, D., Kramer, J., Keppler, D. ubmitted to the EMBL Data Library, July 1 perceiption: The sequence of C. elegans Reference number: Z21123
                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                              Gene: CESP:F45C12.14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Caenorhabditis elegans;Date: 29-Oct-1999 #sequence_revision
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                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                               Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                           Status: preliminary;
                                                                                                                                                                                                                                                                                                                                                                               Cross-references: EMBL:AF016684; PIDN:AAB66212.1;
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                                                                                                                                                                                                                                                                                                                                                                                                              Molecule type: DNA
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ERPGFQIYY 271
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8; Conserv
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7; Conser
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72.7%;
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Pred. No.
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25;
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                                                                                                                                                                                                                                                                                                                                                             2.1; GSPDB:GN00020; CESP:F45C12.14
F45C12
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                                                                                                                                                                                                                                             Length 425;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Masui,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-Oct-1999
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                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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                                                                             hypothetical protein PA3064 [imported] - Pseudomonas aeruginosa (strain PA01) C;Species: Pseudomonas aeruginosa (C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 31-Dec-2000 C;Accession: G83264
                                                                                                                                                                                                              B
                                                                                                                                                                                                                                             Q
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                                                 adman, S.; Yuan,
                                                              R; Stoyer, C.K.; Pham, X.Q.;
                                                                                                                                              G83264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q
                  Nature 406,
                                                                                                                                                                RESULT
                            Lory,
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S

Olson, M.V.

genome

sequence of Pseudomonas aeruginosa PA01, an opportunistic

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Υ.,

Brody,

Erwin, L.L.;

Coulter,

Mizoguchi,

Folger,

S.D.;

Warrener, K.R.; Kas,

Hickey, Larbig,

M.J.;

VQRPQLPLYYS 347

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A;Map position: 5R
C;Superfamily: deubiquinating enzyme
C;Keywords: hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT
S50277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: A; Accession: T38179
                                                                                                                                                  A; Cross-references:
                                                                                                                                                                                                          A; Molecule type: DNA
A; Residués: 1-805 <DIE>
                                                                                                                                                                                                                                                                                                     R;Dietrich,
                                                                                                                                                                                                                                                                                                                                                           A;Title: UBP5 encodes a putative yeast ubiquitin-specific A;Reference number: S50277; MUID:95176708 A;Accession: S50277
                                                                                                                                                                                                                                                                                                                                                                                                                                                          N; Alternate names: protein YEK144c
C; Species: Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Gene: rqs1; SPAC22F3.12c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             submitted to the
                                                                                                                                                                                                                                                                                                                                                                                                                     R; Xiao,
                                                                                                                                                                                                                                                                                                                                                                                                                                   C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Date: 01-Aug-1995 #sequence_revision 01-Sep-1995 #text_change 29-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ubiquitin-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Residues:
                                                                                                                                                                  A; Gene:
                                                                                                                                                                                              A;Cross-references:
                                                                                                                                                                                                                                                                                                                 A;Cross-references: EMBL:U10082;
                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
                                          Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                         east 10,
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Best Local Similarity
Matches 6; Conserv
                               Matches
                                                                                                                                                                                Genetics:
                                                                                                                                                                                                                                                                                                                                  Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      348 ILRKPNLQTYF 358
                                                                                                                                                                                                                                                                                                                                                                                                       W.; Fontanie, T.;
0, 1497-1502, 1994
2 LQRPGLQLYFS 12
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                                                                                                                                                                                                                                                                                                                                  1-805 <XIA>
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                             Similarity 7; Conserv
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er: Z21776
                               Conservative
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                                                                                                                                                                                           EMBL:U18917; NID:g603377; PIDN:AAB64671.1; PID:g603384; MIPS:YER1
                                                                                                                                                 SGD:S0000946; MIPS:YER144c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL: Z54285; NID: ce: strain 972h-;
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                                          60.7%;
63.6%;
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                            Score 37; DB Pred. No. 63; 2; Mismatches
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Pred. No.
                                                                                                                                                                                                                                                                      cerevisiae
                                                                                                                                                                                                                                                                                    December
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                                                         Length 805;
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A;Accession: G83264
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N;Contains: capsid p
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Status: preliminary
                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  rology 155, 77-88, 1986
Title: Cloning full-length dengue type 4 viral DNA sequences:
Reference number: A94352; MUID:87044106
Accession: A94352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Date: 31-Mar-1989 #sequence_revision 31-Mar-1989 #text_change 17-Nov-2000 Accession: A94352; A94364; A26897; A29121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Species: dengue virus type 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cross-references: GB:AE004730; GB:AE004091; NID:g9949154; PIDN:AAG06452.1; GSPDB:GN00:Experimental source: strain PAO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Molecule type: genomic Residues: 774-3386 <MAC>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zhao, B.; Mackow, E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mackow, E.; Makino, Y.; Zhao, B.; Zhang, Y.M.; Markoff, L.; Buckler-White, A.; Guiler cology 159, 217-228, 1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Molecule type: genomic RNA
Residues: 1-776 <ZHA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          litle: The nucleotide sequence of dengue type 4 virus: analysis of genes coding for nucleorence number: A94364; MUID:87293881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ross-references: GB:M17255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          coss-references: GB:M14931
                                                                                                                                                                                                                                                                                                 6-1673/Region: nucleotide-binding motif A (P-loop)
                                                                                                                                                                                                                                                                                                                                                     7-1179/Domain: transmembrane #status predicted <TM6>
5-1343/Product: nonstructural protein NS2a #status predicted <N2A>
1-1473/Product: nonstructural protein NS3b #status predicted <N2B>
-2091/Product: nonstructural protein NS3 #status predicted <NS3>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ords: ATP; capsid protein; envelope protein; glycoprotein; nonstructural protein; 3/Product: capsid protein #status predicted <CAP>
                                                                                                                                                                                                                2486/Product: nonstructural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          79/Product: membrane protein precursor #status predicted 04/Domain: nonterminal signal sequence #status predicted 79/Product: membrane protein #status predicted <MEM>
83/Domain: transmembrane #status predicted <TM3>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         44/Domain: transmembrane #status predicted <TM4>69/Domain: transmembrane #status predicted <TM5>
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                                                                                                                                                3374/Product: nonstructural protein NS4a #status predicted <N4A>
486/Product: nonstructural protein NS4b #status predicted <N4B>
3386/Product: nonstructural protein NS5 #status predicted <NS5>
3386/Product: nonstructural protein NS5 #status predicted <NS5>
6,432,750,903,980,2296,2300,2341,2382,2452,2582,2639,2699,2719,2913,3310/Binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /Product: envelope protein #status predicted <ENV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  yellow fever virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transmembrane #status predicted <TM1>
            Conservative
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pin NS4a; nonstructural protein NS4b
                                                                                                                                                                                                                                                                               DEAH motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nonstructural protein NS1 #status predicted <NS1>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dengue virus type 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Buckler-White, A.; Markoff, L.; Chanock, R.M.; Lai, C.J.; Making
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60.7%;
                                       60.7%; Score 37; 75.0%; Pred. No.
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Pred. No.
         Mismatches
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. 2.4e+02;
0;
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         Indels
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      Gaps.
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Qy 5 PGLQLYFS 12
|||:||:
| Db 466 PGLELYFN 473
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Search completed: June 4, 2001, 12:15:41 Job time: 274 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OM protein - protein search, using sw model
   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                 Score
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61
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YDHE_SCHPO
VGLH_HSV11
VGLH_HSV1E
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FCGI_MOUSE
POLG_BCMVN
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Y824_DEIRA
ATTY_RAT
                                                     RR8_EPIVI
RR8_TOBAC
RR8_MAIZE
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EMBL; X14356; CAA32537.1; EMBL; X14356; CAA32536.1; PIR; S03018; S03018. PIR; S03019; S03019. PIR; A41357; A41357. PIR; B41357; A41357. HSSP; P12319; 1ALT. MIM; 146760; InterPro; IPR003006; Pfam; PF00047; 1g; 3.	"HWW-"http://www.ncbi.nlm.nih.gov/prow/cd/cd64.htm".  "This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).	78-381(1989) BINDS TO TH RECEPTOR. AR LOCATION: PE PRODUCTS: ALTERNATI SCIFICITY: M CCONTAINS		** ** 🗩	LHUMAN STANDARD; PRT; 374 AA.  FCGI_HUMAN STANDARD; PRT; 374 AA.  P12314; P12315;  01-OCT-1989 (Rel. 12, Created) 01-OCT-1989 (Rel. 12, Last sequence update) 01-OCT-2000 (Rel. 40, Last annotation update) 01-OCT-2000 (Rel. 40, Last annotation update) HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR (FC-GAMMA RI) (FCRI) (IGG FC RECEPTOR I) (CD64).

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FCGI_MOUSE

P26151;

P16151;

P26151;

P26152;

P26152
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                                                                "Structure and mapping of the gene encoding movey gamma RI and chromosomal location of the human J. Immunol. 148:1570-1575(1992).
                                                                                                                                                                                         Sears DW. Osman N., Tate B., MCK
Sears DW. Osman N., Tate B., MCK
"Molecular cloning and expression
receptor for IgG.",
J. Immunol. 144:371-378(1990).
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DOMAIN
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
Osman N., Kozak C.A., McKenzie I.F.,
                                                                                                                                                  MEDLINE=92166399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOUSE
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N., Tate B., Mc
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S -> T (IN CAA32536).
2C2AA8103ECF16E6 CRC64;
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HSSP; P12319; 1ALT.
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SEQUENCE
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GENOME POLYPROTEIN [CONTAINS: N-TERMINAL PROTEIN (P1); HELPER
COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); PROTEIN P3; 6 KDA PROTEIN
COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); PROTEIN P3; 6 KDA PROTEIN
[6K1); CYTOPLASMIC INCLUSION PROTEIN (CI); 6 KDA PROTEIN 2 (6K2);
GENOME-LINED PROTEIN (VPG); NUCLEAR INCLUSION PROTEIN A (NI-A) (NIA)
(EC 3.4.22.-) (49 KDA PROTEINASE) (49 KDA-PRO); NUCLEAR INCLUSION
PROTEIN B (NI-B) (NIB) (RNA-DIRECTED RNA POLYMERASE) (EC 2.7.7.48);
COAT PROTEIN (CP)].
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CARBOHYD
Virus Res. 39:13-23(LYYJ).
-!- FUNCTION: HELPER COMPONENT-PROTEINASE IS
TRANSMISSION AND ALSO HAS PROTEOLYTIC AC
                                                                           common mosaic virus (NL3 strain).";
Virus Res. 39:13-23(1995).
                                                                                                                 MEDLINE=96191623; PubMed=8607279;
Fang G.W., Allison R.F., Zambolim
"The complete nucleotide sequence
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InterPro; IPR003006; -.
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European Bioinformatics Institute.
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                                      GLD OR CG1152
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PRINTS; PR00966; NIAPOTYPTASE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute.
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Sutton G.G., Wortman J.R., Y
Brandon R.C., Rogers Y.-H.C.
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Krasney P.A., Carr C.M., Cavener D.R.;
"Evolution of the glucose dehydrogenase
                                                   Cavener D.R., Krasney P.;
"Drosophila glucose dehydrogenase and yea
homologous and share N-terminal homology
                                                                                                                                                                                                                                                                                                                                                         SIMILARITY TO YEAST ALCOHOL OXIDASE MEDLINE=91163320; PubMed=2002763;
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"Molecular structure and transformation
gene in Drosophila melanogaster.";
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Beeson K.Y., Benos P.V
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Amanatides P.G., Scherer S.
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                                                                                                                                                                Perlaky S.,
                                                                                                                                                                                         SELENOCYSTEINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Science
                                                                                                                                                                                                                                                                                                                                                                                                                                            Genetics 120:475-484(1988)
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COFACTOR: FAD
                         REDUCED ACCEPTOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nome sequence of Dro
287:2185-2195(2000)
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                                                                                                                                                                                                                                                Evol.
                                                                                                                                                                Merritt K.,
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                                                                                                                                                                                                                                                8:144-150(1991).
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V., Berman B.P., Bhandari D., Bolshakov S
                                                                                                                                                                Cavener D.;
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nation of the glucose dehydro
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ogy with other
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                                                                                                                                      of Gld.";
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Smith H.O.,
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Bioinformatics Institute.

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SE_CYS
CONFLICT
SEQUENCE
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                                                                                     TISSUB-Embryonic lung;
MEDLINE-97064168; PubMed-8906787;
Stone D.M., Hynes M., Armanini M.
Scott M.P., Pennica D., Goddard A.
de Sauvage F., Rosenthal A.;
                                                                                                                                                                  Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                30-MAY-2000 (Rel.
30-MAY-2000 (Rel.
01-OCT-2000 (Rel.
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Xie J., Murone M., L
Lam C.-W., Hynes M.,
de Sauvage F.J.;
                             SEQUENCE FROM N.A., AND VARIANTS BCC LEU-535 AND GLN-562 MEDLINE=98082974; PubMed=9422511;
                                                       Sonic hedgehog.";
Nature 384:129-134(1996).
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S06628; S06628.
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39, Last sequence update)
40, Last annotation updat
PRECURSOR (SMO).
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        uon s.-M., Rÿan A.,
Goddard A., Rosent
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77.8%;
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           Rosenthal
                                                                                             Swanson T.A.,
Phillips H.,
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         Gu 'Q.,
chal A.,
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                                                                             candidate receptor for
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         Zhang C., Bonifas J.M., Epstein E.H. Jr.,
                                                                                                                                                                                                                                                                                                                                                               Length 625
                                                                                                                                                                                                                                                                                                                                                                                          CRC64;
                                                                                              Gu Q., Jo
Noll M.,
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                                                                                                        Johnson R.L.,
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RESULT 6
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CARBOHYD
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EMBL; AF114819; AAD17202.1;
EMBL; AF114820; AAD17202.1;
MIM; 601500;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
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-i- FUNCTION: G PROTEIN-COUPLED RECEPTOR THAT PROBABLY ASSOCIATES WITTE PATCHED PROTEIN (PTCH) TO TRANSDUCE THE HEDGEHOG'S PROTEINS SIGNAL BINDING OF SONIC HEDGEHOG (SHH) TO ITS, RECEPTOR PATCHED THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED
                                                       SEQUENCE
                                                                              VARIANT
                                                                                                                                                                                                                                                                                                                              SIGNAL
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Developmental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Activating Smoothened mutations in sporadic basal-cell carcinoma.";
                                                                                                                                                                                                                                                                                                                  CHAIN
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l Similarity
8; Conservat
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           62.3%;
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 Score 38; DB Pred. No. 19; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                   Transmembrane; Glycoprotein;
                                                                                                                                                                                                                                    3 POTENTIAL.
CYTOPLASMIC (POTENTIAL).
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2 POTENTIAL.
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N-LINKED (GLCNAC...)
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CYTOPLASMIC (POTENTIAL)
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CYTOPLASMIC (POTENTIAL):
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                                                                  /FTId=VAR_007848.

R -> Q (IN BCC; SOMATIC MUTATION)
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                                                                                                    -> L (IN BCC; SOMATIC
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                                                         8B4C459B34D13F83 CRC64;
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IN BASAL CELL CARCINOMAS (BCC),
         DB
19;
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"Cloning of a mouse Smoothened cDNA and expression patterns of hedgehog signalling molecules during chondrogenesis and cartilage differentiation in clonal mouse EC cells, ATDC5.";

Biochem. Biophys. Res. Commun. 235:142-147(1997).

"I-FUNCTION: G PROTEIN (PTCH) TO TRANSDUCE THE HEDGEHOG'S PROTEINS SIGNAL. BINDING OF SONIC HEDGEHOG (SHH) TO ITS RECEPTOR PATCHED IS THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED.
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Mammalia; Eutheria;
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-!- SIMILARITY: BELONGS TO FAMILY FZ/SMO OF G-PROTEIN COUPLED
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Query Match Best Local Matches

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Scott M.P., Pennica D.,
de Sauvage F., Rosenthal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         between the Swiss Institute of Bioinf
the European Bioinformatics Institute.
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SIMILARITY: BELONGS TO FAMILY FZ/SMO OF G-PROTEIN COUPLED
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THE PATCHED PROTEIN: (PTCH) TO TRANSDUCE THE HEDGEHOG'S PROTEINS
SIGNAL. BINDING OF SONIC HEDGEHOG (SHH) TO ITS RECEPTOR PATCHED
THOUGHT TO PREVENT NORMAL INHIBITION BY PATCHED OF SMOOTHENED
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                                                                                                                                                                                                                                                                                                                                               PF01534; Frizzled; PF01392; Fz; 1.
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                                                                                                                                                                                                                                                                                                      protein
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ca D., Goddard A., Phillips H., Noll M.,
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Rodentia;
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87375
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IN EMBRYO, FOUND IN THE EARLY NEURAL
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N-LINKED (GLCNAC...) (P
N-LINKED (GLCNAC...) (P
, 27B88178B2565FB9 CRC64;
                                                                                                                                4 POTENTIAL.

EXTRACELLULAR (POTENTIAL).

5 POTENTIAL.
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P18479; Q8
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-!- FUNCTION: CYTOPLASMIC INCLUSION PROTEIN HAS HELICASE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-95146958; PubMed-7844540; Wisler G.C., Purcifull D.E., Hiebert "Characterization of the P1 protein yellow mosaic virus.";
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15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GENOME POLYPROTEIN (CONTAINS: N-TERMINAL PROTEIN (P1);
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PTM: THE VIRAL RNA OF POTYVINGUSES IS EXPRESSED AS A SINGLE
POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTECLYTIC
PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT
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SIMILARITY: BELONGS TO THE POTYVIRUSES POLYPROTEIN FAMILY SIMILARITY: HC PROTEINASE BELONGS TO PEPTIDASE FAMILY C6. SIMILARITY: NI-A PROTEINASE, BELONGS TO PEPTIDASE FAMILY C4.

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EMBL; L31350; AAA65559.1; -. EMBL; M35095; AAA48511.1; -.
InterPro;
             InterPro;
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                                                                     JH0103;
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             IPR001650;
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Piam; Piam;

Pfam; InterPro;

PF00863; Peptidase\_C4;
PF00851; Peptidase\_C6; PF01577; Poty\_P1;

IPR002540;

Pram;

PF00767;

Poty\_

coat;

RNA\_dep\_RNA\_pol;
helicase\_C; 1.

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Matches 5
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15-JUL-1999 (Rel. 38, Last sequence update)
15-JUL-1999 (Rel. 38, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation update)
GENOME POLYPROTEIN [CONTAINS: N-TERMINAL PROTEIN (P1); HELPER
COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); PROTEIN P3; 6 KDA PRO
1 (6K1); CYTOPLASMIC INCLUSION PROTEIN (CI); 6 KDA PROTEIN 2 (6K2)
GENOME-LINKED PROTEIN (VPG); NUCLEAR INCLUSION PROPEIN A (NI-A) (I

(EC 3.4.22.-) (49 KDA PROTEINASE) (49 KDA-PRO); NUCLEAR INCLUSION
                                                                              This
                                                                                                                                                                                                                                                                               Lee K.C., Wong S.M.;
Submitted (JUL-1997)
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                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                      POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTEOLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT INDIVIDUAL PROTEINS.
SIMILARITY: HC PROTEINASE BELONGS TO PEPTIDASE FAMILY C6.
SIMILARITY: NI-A PROTEINASE BELONGS TO PEPTIDASE FAMILY C6.
SIMILARITY: BELONGS TO THE POTYVIRUSES POLYPROTEIN FAMILY
                                                                                                                                                                                                                               FUNCTION: HELPER COMPONENT-PROTEINASE IS REQUIRED TRANSMISSION AND ALSO HAS PROTECLYTIC ACTIVITY. FUNCTION: CYTOPLASMIC INCLUSION PROTEIN HAS HELICA MAY BE INVOLVED IN REPLICATION.
                                                                                                                                                                                       FUNCTION: NUCLEAR INCLUSION PROTEIN A HAS PROTEOLYTIC ACTIVITY. PTM: VPG IS COVALENTLY LINKED TO THE GENOMIC RNA. PTM: THE VIRAL RNA OF POTYVIRUSES IS EXPRESSED AS A SINGLE
                                                                SWISS-PROT entry is copyright. It is produced through a collaboration sen the Swiss Institute of Bioinformatics and the EMBL outstation -
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6 KDA PROTEIN 2.
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G -> S (IN REF. 2).
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71;
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15-DEC-1998 (Rel.
15-DEC-1998 (Rel.
15-DUL-1999 (Rel.
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15-DEC-1998 (Rel. 37, Last sequence update)
15-UL-1999 (Rel. 38, Last sequence update)
GENOME POLYPROTEIN [CONTAINS: N-TERMINAL PROTEIN (P1); HELPER
COMPONENT PROTEINASE (EC 3.4.22.-) (HC-PRO); PROTEIN P3; 6 KDA PROTI
1 (6K1); CYTOPLASMIC INCLUSION PROTEIN (CI); 6 KDA PROTEIN 2 (6K2);
GENOME-LINKED PROTEIN (VPG); NUCLEAR INCLUSION PROTEIN A (NI-A) (NII)
(EC 3.4.22.-) (49 KDA PROTEINASE) (49 KDA-PRO); NUCLEAR INCLUSION
PROTEIN B (NI-B) (NIB) (RNA-DIRECTED RNA POLYMERASE) (EC 2.7.7.48);
COAT PROTEIN (CP)].
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Pfam; PF00767; Poty_coat; 1.
Pfam; PF00680; RNA_dep_RNA_pol; 1.
Pfam; PF00680; RNA_dep_RNA_pol; 1.
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Hydrolase; Transferase; Thiol protease; RNA-directed
Hydrolase; Transferase; Covalent protein-RNA link;
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                                                                                         "Characterization of the P1 protein yellow mosaic virus.";
J. Gen. Virol. 76:37-45(1995).
                                                                                                                                              SEQUENCE FROM N.A. MEDLINE-95146958; PubMed-7844540;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         1972 IINKPGIQAYF 1982
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                                                                                                                                 Wisler G.C., Purcifull D.E., Hiebert E.;
                                                                                                                                                                                                               Viruses; ssRNA positive-strand viruses, no DNA stage;
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FUNCTION: HELPER COMPONENT-PROTEINASE IS REQUIRED FOR APHID TRANSMISSION AND ALSO HAS PROTEOLYTIC ACTIVITY. FUNCTION: CYTOPLASMIC INCLUSION PROTEIN HAS HELICASE ACTIVITY. FUNCTION: EXPOLED IN REPLICATION.

FUNCTION: NUCLEAR INCLUSION PROTEIN A HAS PROTEOLYTIC ACTIVITY. FUNCTION: NUCLEAR INCLUSION PROTEIN A HAS PROTEOLYTIC ACTIVITY. PTM: VPG IS COVALENTLY LINKED FOO THE GENOMIC RNA.

PTM: THE VIRAL RNA OF POTYVIRUSES IS EXPRESSED AS A SINGLE
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                                                                                                                   and coding region
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Helicase;
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01-NOV-1997
30-MAY-2000
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InterPro; IPRO0:
Pfam; PF00863; I
Pfam; PF00851; I
Pfam; PF001577; I
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SEQUENCE
SEQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-97426617; PubMed-9278503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRINTS; PR00966; NIAPOTYPTASE.
Hydrolase; Transferase; Thiol protease; RNA-directed RNA polymerase;
Coat protein; Polyprotein; Covalent protein-RNA linkage; Helicase;
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                                                NCBI_TaxID=562;
                                                            Bacteria; P:
Escherichia
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PF00680; RNA_dep_RNA_pol;
PF00271; helicase_C; 1.
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                                                                       Proteobacteria;
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7; Poty_P1; 1.
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(Rel. 35, Last sequence update)
(Rel. 39, Last annotation update)
TRANSCRIPTIONAL REGULATOR IN DMSC-PFLA
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55E51B455C20C537 CRC64;
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. 72;
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TO PEPTIDASE FAMILY C4
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01-NOV-1995 (Rel. 3
01-OCT-2000 (Rel. 4
HYPOTHETICAL 54.4 K
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01-NOV-1995
01-OCT-2000
                         Lye G., Churcher C.M., Barrell B.G., Rajandres Submitted (SEP-1995) to the EMBL/GenBank/DDBJ-!- SIMILARITY: CONTAINS 1 RGS DOMAIN.
                                                                                                                                                   Schizosaccharomyces pombe (Fission yeast).
Eukaryota; Fungi; Ascomycota; Schizosaccharomycetales; Schizosaccharomyc
This SWISS-PROT entry is copyright.
                                                                                                                           Schizosaccharomyces.
NCBI_TaxID=4896;
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                                                                                      STRAIN-972
                                                                                                 SEQUENCE FROM N.A.
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Q09777;
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A 718-kb DNA sequence of the Escherichia coli K-12 genome corresponding to the 12.7-28.0 min region on the linkage map. DNA Res. 3:137-155(1996).

-I- SIMILARITY: BELONGS TO THE LYSR FAMILY OF TRANSCRIPTIONAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oshima T., Aiba H., Baba T., Fujita K., Hayashi K., Honjo A., Ikemoto K., Inada T., Itoh T., Kajihara M., Kanai K., Kashimoto I Kimura S., Kitagawa M., Makimo K., Masuda S., Miki T., Mizobuchi Mori H., Motomura K., Nakamura Y., Nashimoto H., Nishio Y., Saita
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EcoGene; EG13698; ycaN.
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PS00046; HTH LYSR FAMILY; 1.

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PS00046; DNA-binding.

B-7-H MOTIF (POTENTIAL).

302 AA; 34294 MW; 8FFF02FB02B24B73 CRC64;
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Pred. No.
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C., Yamamoto
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DEDS_YEAST

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P39944;

T 01-FEB-1995 (Rel. 31, Created)

T 01-FEB-1995 (Rel. 31, Last sequence update)

DT 01-NOV-1995 (Rel. 32, Last annotation update)

DE UBIQUITIN CARBOXYL-TERMINAL HYDROLASE 5 (EC 3.1.2.1

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C STRAIN-5288C / AB972;

A Dietrich F.S., Mulligan J.T., Hennessey K.M., Allen E., Araujo R.,

A Aviles.E., Berno A., Brennañ T., Carpenter J., Chen E., Cherry J.M.

Chung E., Duncan M., Guzman E., Hartzell G., Hunicke-Smith S.,

A Hyman R., Kayser A., Komp C., Lashkari D., Lew H., Lin D.,

Mosedale D., Nakahara K., Namath A., Worgren R., Oefner P., Oh C.,

A Mosedale D., Nakahara K., Namath A., Worgren R., Oefner P., Oh C.,

A Petel F.X., Roberts D., Sehl P., Schramm S., Shogren T., Smith V.,

A Taylor P., Wei Y., Yelton M., Botstein D., Davis R.W.;

Submitted (DEC-1994) to the EMBL/GenBank/DDBJ databases.

-!- CATALYTIC ACTIVITY: UBIQUITIN C-TERMINAL THIOLESTER + H(2)O =

UBIQUITIN + A THIOL.

C -!- SIMILARITY: BELONGS TO PEPTIDASE FAMILY C19; ALSO KNOWN AS

FAMILY 2 OF UBIQUITIN CARBOXYL-TERMINAL HYDROLASES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=95176708; PubMed=7871889;
Xiao W., Fontanie T., Tang M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Saccharomyces cerevisiae (Baker's yeast)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Fungi;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomycetales;
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"In vitro processing of dengue virus st
the pre-membrane protein.",
J. Virol. 63:3345-3352(1989),
-i- FUNCTION: THE SMALL PROTEINS NS2A,
HYDROPHOBIC, SUGGESTING A POSSIBLE
                                                                                                                                                          SEQUENCE OF 774-3386 FROM N.A.
SEQUENCE OF 774-3386 FROM N.A.
MEDLINE-87293881; PubMed-3039728;
Mackow E., Makino Y., Zhao B., Zhang Y.M., Markoff L.,
Mackow E., Makino Y., Zhao B., Zhang Y.M., Markoff L.,
Buckler-White A., Guiler M., Chanock R., Lai C.J.;
"The nucleotide sequence of dengue type 4 virus: analysis of genes coding for nonstructural proteins.";
Virology 159:217-228(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAR-1989 (Rel. 10, Created)
01-JUL-1989 (Rel. 11, Last sequence update)
30-MAY-2000 (Rel. 3), Last annotation update)
30-MAY-2000 (Rel. 39, Last annotation update)
GENOME POLYPROTEIN [CONTAINS: CAPSID PROTEIN C (CORE PROT PROTEIN (ENVELOPE GLYCOPROTEIN M); MAJOR ENVELOPE PROTEIN NONSTRUCTURAL PROTEINS NS1, NS2, NS4A AND NS4B; HELICASE RNA-DIRECTED RNA POLYMERASE (EC 2.7.7.48) (NS5)]
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e; Nonstructural protein
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Plavi_NS4B; 1
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lavi_helicase; 1.
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MEDLINE-95350630; PubMed-7542800;

Fleischmann R.D., Adams M.D., White O., Clayton R.A., Kirkness E.F.

Kerlavage A.R., Bult C.J., Tomb J.-F., Dougherty B.A., Merrick J.M.

MCKenney K., Sutton G., Fitzhugh W., Fields C.A., Gocayne J.D.,

Scott J.D., Shirley R., Liu L.-I., Glodek A., Kelley J.M.,

Weidman J.F., Phillips C.A., Spriggs T., Hedblom E., Cotton M.D.,

Utterback T.R., Hanna M.C., Nguyen D.T., Saudek D.M., Brandon R.C.,

Fine L.D., Fritchman J.L., Fuhrmann J.L., Geoghagen N.S.M.,

Gnehm C.L., McDonald L.A., Small K.V., Fraser C.M., Smith H.O.,
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PROSITE; PS00402; BPD_TRANSP_INN_MEMBR; 1.
Transport; Phosphate transport; Transmembrane;
DOMAIN 1 22 CYTOPLASMIC (POTEN
TRANSMEM 23 43 POTENTIAL.
                                                                                                               modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no restructed by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                    Science 269:496-512(1995).

-i- FUNCTION: PART OF THE BINDING-PROTEIN-DEPENDENT TRANSPORT-
-i- FUNCTION: PROBABLY RESPONSIBLE FOR THE TRANSLOCATION
SUBSTRATE ACROSS THE MEMBRANE (BY SIMILARITY).

-i- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. INNER MEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1995
01-NOV-1995
                                                                           EMBL; U32818; AAC23027.1; TIGR; HI1382; -
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Bacteria; Proteobacteria;
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                                                    Pfam; PF00528; BPD_transp; 1.
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ALIGNMENTS

## RESULT Q92663 Query Match Best Local Similarity Matches 12; Conser EMBL; L03418; AAA36049.1; HSSP; P12319; TALT: INTERPRO; IPRO3006; -. PFAM; PF00047; 19; 3. PRODOM; PD002534; -; 1. Q92663; 01-FEB-1997 (TrEMBLrel. 01-FEB-1997 (TrEMBLrel. 01-OCT-2000 (TrEMBLrel. SEQUENCE "Novel Fc gamma receptor I family gene products in human mononuclear cells.", MEDLINE-93055454; PubMed-1430234; Porges A.J., Redecha P.B., Doebele R., Kimberly R.P.; Eukaryota; Metazoa; Mammalia; Eutheria; 01-OCT-2000 (TrEMBLrel. FC GAMMA RECEPTOR I. Q92663 TISSUE-BLOOD, SEQUENCE FROM N.A. iomo sapiens (Human) 7. Clin. Invest. 90:2102-2109(1992). CBI\_TaxID=9606, 100.0%; Score 61; DB 4; ilarity 100.0%; Pred. No. 0.0013; Conservative 0; Mismatches C 374 AA; PRELIMINARY; Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo. 42632 MW; 15, Created) Last sequence update) Last annotation update) D33D59398CEEA699 CRC64; PRT; 374 Pan L.C., B Length 374; Salmon J.E., Indels Ö Gaps

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01-OCT-2000 (TrEMBLrel. 1
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FC GAMMA RECEPTOR I.
              Yan Y., Li X., Wang A., Zhang G.;
"Molecular cloning and identification of full-length cDNA high affinity Fc receptor for bovine IgG (Fc gamma RI).";
Vet. Immunol. Immunopathol. 75:151-159(2000).
EMBL; AF162866; AAF80477.1;
                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea Bovidae; Bovinae; Bos. NCBI_TaxID=9913;
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PubMed=1402657;
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Catarrhini; Hominidae;
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                                    Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Eterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
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MEDLINE=20196006;
        SEQUENCE FROM N.A. STRAIN-BERKELEY;
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SEQUENCE 612 AA; 66794 MW;
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EMBL; U63325; AAB48020.1; -
EMBL; U63324; AAB48020.1; JOINED.
EMBL; U63324; AAB48020.1; JOINED.
FLYBASE; FB9n0013875; Dsim\Gld.
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Eukaryota; Metazoa; Arthropoda; Tracheata; He
Eukaryota; Neoptera; Endopterygota; Diptera;
Ephydroidea; Drosophilidae; Drosophila.
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Hamblin M.T., Aquadro C.F.;
"High nucleotide sequence variati
in Drosophila simulans is consist
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Best Local Similarity //..
Thes 7; Conservative
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Q9UZX2;
01-MAY-2000
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                                                                                                                                                                                                                               Pyrococcus abyssi.
Archaea; Euryarchaeota;
NCBI_TaxID=29292;
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01-JUN-2000
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                                                                                                                                              SEQUENCE FROM N.A
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         Pyrococcus abyssi genome
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G.G., Scherer S.E., I
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77.88;
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sequence: insights into archaeal
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Last annotation update)
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Pred.
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1.
BC162CDE4252BCE0 CRC64;
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Cadieu E., Center A., Chandra
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.A., Galle
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         chromosome
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RESULT Q9KFN1

Q9KFN1;

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RESULT
Q05912
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Submitted (JUL-1999) to the ENBL; AJ248286; CAB49934.1; -
Hypothetical protein.
SEQUENCE 393 AA; 44932 MW;
                                                                                                                                                                                                                     PFAM;
                                                                                                                                                   Polyprotein; Covalent protein-RNA NON_TER 1 1
                                                                                                                                                                                                                                                     INTERPRO;
                                                                                                                                                                                                                                                               INTERPRO;
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"Nucleotide sequence of the 3' half of zucchini yellow mosaic virus
(Singapore isolate) genome encoding the 4K protein, protease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zucchini yellow mosaic virus (strain Si
Viruses; ssRNA positive-strand viruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENOME POLYPROTEIN (FRAGMENT)
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01-OCT-2000
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                                                                                                                         SEQUENCE
                                                                                                                                                                              Hydrolase;
                                                                                                                                                                                             PRINTS; PR00966; NIAPOTYPTASE
                                                                                                                                                                                                                                      INTERPRO;
                                                                                                                                                                                                                                                                                          - ! - SIMILARITY: TO OTHER POTYVIRUSES POLYPROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic Acids Res. 21:1317-1317(1993).
-!- FUNCTION: CYTOPLASMIC INCLUSION PROTEIN HAS HELICASE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=93219099; PubMed=8464715;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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179 IINKPGIQAYF 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                 olymerase and coat protein."
                                                                Local
                           1 LLQRPGLQLYF 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 LLQRPGLQLYF
                                                                                                                                                                                                                                                                                                                    PTM: THE VIRAL RNA OF POTYVIRUSES IS EXPRESSED AS A SINGLE POLYPROTEIN WHICH UNDERGOES POSTTRANSLATIONAL PROTEOLYTIC PROCESSING RESULTING IN THE PRODUCTION OF AT LEAST EIGHT
                                                                                                                                                                                                                                                                                                                                                            IT, VPG, IS COVALENTLY LINKED PROTEOLYTIC ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION: NUCLEAR INCLUSION PROTEIN A, OR THE N-TERMINAL PART
                                                                                                                                                                                                                                                                                                                                                                                                      MAY BE INVOLVED IN REPLICATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LMTKPGIEVYF 199
                                                                                                                                                                                                                                                                                                       NDIVIDUAL PROTEINS
                                                                                                                                                                                                          PF00680; RNA_dep_RNA_pol;
PF00863; Peptidase_C4; 1.
                                                                                                                                                                                                                                                                             X68509; CAA48521.1;
                                                     Similarity
5; Conserv
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5; Conser
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IPR001730;
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                                                                                                                                                                Protease; RNA-directed RNA polymerase; Coat protein; n; Covalent protein-RNA linkage.
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1016
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                                                     Conservative
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                                                                62.3%;
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01, Last sequence 15, Last annotation
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Last annotation update)
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                                                                   Score 38;
Pred. No.
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nd viruses, no DNA stage; Potyviridae;
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                                                  Mismatches
                                                                                                                         6887BEAACB24E387 CRC64;
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Best Local
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Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson /
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.
Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen
Smaldon N., Smith A., Sonnhammer K., Saunders D., Shownkeen
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01-JAN-1998 (TrEMBLrel.
01-JAN-1998 (TrEMBLrel.
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ABC TRANSPORTER (
 Submitted (JUL-1997) to the EMBL; AF016684; AAB66212.1; INTERPRO; IPR001810; -. PFAM; PF00646; F-box; 1. SEQUENCE 425 AA; 48488 MW
                                                                                                                                     [2]
SEQUENCE FROM N.A..
                                                                                                                                                                                                                                                                                                                                                                                                 Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda;
Rhabditidae; Peloderinae; Caer
                                                                                                              Johnson D., Kramer J.,
Submitted (AUG-1997) to
                                                                                                                                                                                                            Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Watten Matson A., Weinstock L., Wilkinson Sproat J., Wohldman
                                                                                                                                                                                                                                                                                                                                  STRAIN=BRISTOL N2;
MEDLINE=94150718; PubMed=7906398;
                                                                 Waterston
                                                                          SEQUENCE FROM N.A.
STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                016743
                                                                                                                                                                          Nature 368:32-38(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Takami H., Nakasone K., Taka
Submitted (MAR-2000) to the
EMBL; AP001508; BAB04167.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillus/Staphylococcus group; Bacillus. NCBI_TaxID=86665;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
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(TrEMBLrel. 15,
(TrEMBLrel. 15,
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  48488 MW;
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63.6%;
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the EMBL/GenBank/DDBJ databases
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Caenorhabditis.
                                                EMBL/GenBank/DDBJ
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Last annotation updat
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Pred.
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D42932E3A29F6A7D CRC64;
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No.
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                                                databases
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Q9K451
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Best Local S
Matches 8
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Best Local
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01-MAY-2000
01-JUN-2000
                                                  Q9VY94;
STRAIN-BERKELEY;
MEDLINE=20196006; PubMed=10731132;
Adams M.D., Celniker S.E., Holt R.A.,
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01-OCT-2000 (TrEMBLrel. 15, Creat
01-OCT-2000 (TrEMBLrel. 15, Last
01-OCT-2000 (TrEMBLrel. 15, Last
CYTOCHROME C OXIDASE POLYPEPTIDE
                                                                                                    GRIP91.
                                                                                                              GRIP91 PROTEIN.
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EMBL; AL359215; CAB94657.1; -.
SEQUENCE 573 AA; 64041 MW; 1F
                               SEQUENCE FROM N.A.
                                                                              Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda;
                                                                                                                                                                                                                                                                                                                                                     Redenbach M., Kieser H.M., Denapaite D., Kinashi H., Hopwood D.A.,
                                                                                                                                                                                                                                                                                                                                                                         STRAIN-A3(2);
                                                                                                                                                                                                                                                                                                                                                                                                       Submitted
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Thomson N.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=A3(2);
Seeger K.J.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Streptomyces coelicolor.
Bacteria; Firmicutes; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; St
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      (JUN-2000)
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                                                                                                                      (TrEMBLrel. 13, (TrEMBLrel. 13, (TrEMBLrel. 14,
                                                                                                                                                             PRELIMINARY;
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77.8%;
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Pred. No. 53;
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r A3(2) ch
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73;
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53;
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                                                                     Brachycera; Muscomorpha;
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                                                                                                                                                                                                                                                                      Length 573
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Evans

C.A.,

Gocayne J.D.,

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Best Local
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Q9W1Q7;
01-MAY-2000
01-MAY-2000
01-JUN-2000
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                                                                                                                                              Drosophila melanogaster (Fruit fly),
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda;
Pterygota; Neoptera; Endopterygota; Diptera; Brachyce
Ephydroidea; Drosophilldae; Drosophila.
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MEDLINE-20196006; PubMed-10731132; Adams M.D., Celniker S.E., Holt R. Amanatides P.G., Scherer S.E., Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C.,
Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleisc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S
Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Brandon R.C.,
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Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
                                                                   STRAIN-BERKELEY;
                                                                                     SEQUENCE FROM N.A.
                                                                                                                                NCBI_TaxID=7227;
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917 AA; 103706 MW; 6AEE88C211D256BB
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Doyle C., I
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Last sequence update)
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Evans C.A.,
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Gocayne J.A., Galle
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RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guán P., Harris M.,
RA Harris N.L., Harvey D., "Heinan T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., "Heinan T.J., Wei M.-H., Ibegwam C.,
RA Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
Liasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA McIulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,
RA Reinert K., Remington K., Sannders R.D.C., Scheeler F., Shen H.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,
RA Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
Walliams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q. C.,
Ran The Resource of Drospothia melanogaster "."
The Resource of Drospothia melanogaster "."
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Q55990;
Q1-NOV-1996
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Sugiura M., Tabata S.;
"Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. I. Sequence features in the 1 Mb region from map positions 64% to 92% of the genome.";
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Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshak
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                                                                                                                                                                   MEDLINE=96127529; PubMed=8590279; Kaneko T., Tanaka A., Sato S., Ko
                                                                                                                                                                                                                                                     SEQUENCE FROM N.A. STRAIN-PCC6803;
                                                                                                                                                                                                                                                                                                                                                                                                        Synechocystis sp. (strain Bacteria; (
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nce 287:2185-2195(2000).
; AE003460; AE446996.1; -.
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64 AA; 7304 MW; 0A
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Rogers Y.-H.C.,
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Baldwin D.,
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SEQUENCE FROM N.A. MEDLINE=97061201;

Sato S.,

PubMed=8905231;

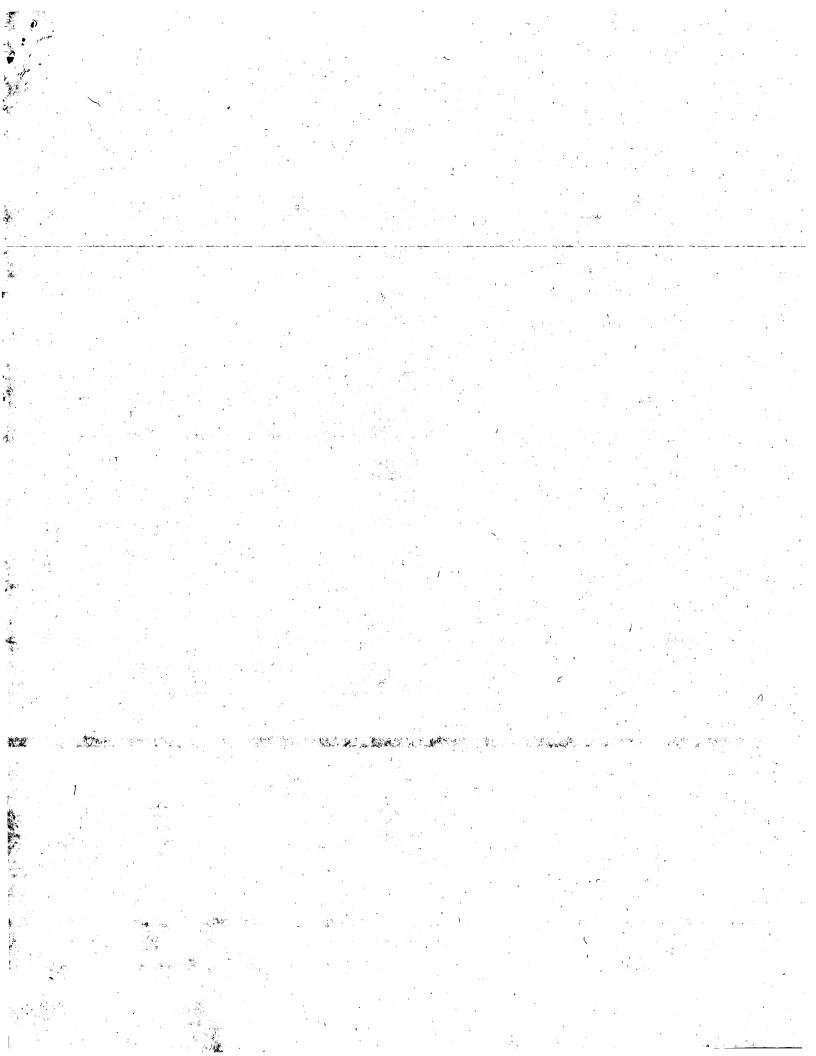
Tanaka A.,

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Best Local Similarity 58.3%;
Matches 7; Conservative
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Best Local Similarity 63.6
Q9Y477 PRELIMINARY;
Q9Y477;
01-NOV-1999 (TIEMBLIFEL 12, C
01-NOV-1999 (TIEMBLIFEL 12, L
01-NOV-1999 (TIEMBLIFEL 12, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Q69075 PRELIMINARY;
Q69075;
Q1-NOV-1996 (TrEMBLrel. 01, C
Q1-NOV-1996 (TrEMBLrel. 01, L
Q1-NOV-1998 (TREMBLREL. 08, L
GLYCOPROTEIN H (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Excretion of non-infectious virus particles lacking glycoprotein H by a temperature-sensitive mutant of herpes simplex virus type 1: evidence that gH is essential for virion infectivity.";
J. Gen. Virol. 69:1147-1156(1988).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human herpesvirus 1.
Viruses; dsDNA viruses, no RNA stage;
Alphaherpesvirinae; Simplexvirus.
NCBI_TaxID=10298;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synechocystis sp. strain PCC6803. entire genome and assignment of pDNA Res. 3:109-136(1996). BAA10772.1;
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Miyajima N., Hirosawa M., Sugiura M.,
Hosouchi T., Matsuno A., Muraki A., N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Desai P.J.,
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                                                                                                                                                                                                                                                 123 LLRSPGRYVYFS 134
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tabata S.
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||: | || ||
94 LLESPSLQKYF 104
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166 AA;
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17805 MW;
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63.6%;
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                                                                                                                171 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Herpesviridae;
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30;
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                                  Query Match
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Matches 7
                                                                         EMBL; AL022575; CAA18605.1; -.
NON_TER 1 1 1
NON_TER 171 171
SEQUENCE 171 171
                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                        DJ159A15.1 (FRAGMENT).
DJ159A15.1.
                                                                                                                                               NCBI_TaxID=9606;
109 LLKRPQLELY 118
                  1 LLQRPGLQLY 10
                                   Similarity 70.0
7; Conservative
                                          59.0%;
70.0%;
                                  Score 36; DB Pred. No. 32; 2; Mismatches
                                    2,
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32;
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Search completed: Job time: 587 sec June 4, 2001, 12:23:24



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Title:
Perfect score:
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protein search, using sw model
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length: 2000000000
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Match
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Gapop.10.0 , Gapext 0.5
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R20811
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Y33183
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Human FCRI (CDNA C
Human FCRI (CDNA C
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                                                                                                                            Human macrophage-s
Human FcRI (cDNA c
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54.7	54.7	• •	•		•	56.2		57.8	57.8	57.8	57.8	57.8	57.8	57.8	57.8	•	57.8	•	•	57.8	•		•	•	57.8	•	•	62.5	•	100.0	100.0	100.0	1000
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W85603	B21429	B21433	B21431	G14207	G14208	G14209	R54218	R66462	R38470	в29019	R92705	R44514	G38488	G37164	G28965	G07160 ·	G38489	G37165	G28966	G07161	G38490	G37166	G28967	G07162	B01787	R07302	Y87523	R12428	W86195	в43683	Y96226	Y96134	70
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saminid	Cone snail alpha-	Cone snail alpha-	_	Arabidopsis thalia	Arabidopsis thalia	Arabidopsis thalia	'L. lactis branched	ALL-1 (acute lymph	ALL-1 protein.	O.	MLL cDNA clone	MLL amino acid	Arabidopsis thalia			Arabidopsis thal	Arabidopsis tha	Arabidopsis tha	Arabidopsis thal	Arabidopsis thalia		Arabidopsis thalia		Arabidopsis thalia	Escherichia coli	C-terminal sequenc	Conotoxin peptide	Hybrid Fc(gamma)RI	Human Fc receptor	Human cancer assoc	Human high affinit	Human macrophage-	2

## ALIGNMENTS

RESULT W60556

Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; detecell transformation; mutation; anti; oncogene. WPI; 1998-240964/21. De Kruif CA, 08-OCT-1996; 07-OCT-1997; WO9815833-A1. Synthetic. Oligopeptide from extracellular domain of CD64. 18-AUG-1998 W60556; W60556 standard; peptide; 12 (UYUT-) RIJKSUNIV UTRECHT. 16.-APR-1998 (first entry) Logtenberg T; 96EP-0202791 97WO-NL00557. A A analysis; detection;

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies

e.g. identifying

Example 1; Page

29; 40pp; English

W60537-61

are derived from the

extracellular

Sequence by mutation

12 AA;

in (anti)oncogenes.

cc adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a comethod for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the cc surface of a replicable display package, synthesising oliqopeptides comprises displaying the peptide on the cc surface of a replicable display package, synthesising oliqopeptides comprises derived from the target protein on a solid phase, and testing for binding between the peptide and oliqopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two cepitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of immunofluorescent analysis, and also to detect cell transformation caused by mutation in (antipocarenes)

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Query Match
Best Local Similarity
Watches 12; Conserv
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                                       Maxwell KF,
WPI; 1999-539978/45
                          Powell MS;
                                                     Baell JB,
                                                                                                        11-SEP-1998;
06-FEB-1998;
                                                                                                                                                04-FEB-1999;
                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                       anti-inflammatory;
                                                                                                                                                                                                                                                                                             drug design; therapy; inhibitor;
                                                                                                                                                                                                                                                                                                                      Fc receptor;
                                                                                                                                                                                                                                                                                                                                               Human sFc-gammaRI protein fragment.
                                                                                                                                                                                                                                                                                                                                                                                                       Y33183
                                                                                                                                                                                                                                                                                                                                                                                                                             Y33183 standard; Protein;
                                                                              (ILEX-) ILEXUS PTY LTD.
                                                                                                                                                                          12-AUG-1999
                                                                                                                                                                                                  WO9940117-A1
                                                                                                                                                                                                                                                                                  tissue damage;
                                                                                                                                                                                                                                                                                                            atomic
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                                                                                                                                                                                                                                                                  tissue damage; hypersensitivity; infinifiammatory modulator; Fc-gammaRIa;
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                                                                                                                                                                                                                                                                                                       coordinate; bloactive compound design; computer-assisted
                                     Epa V, Gar
7, McCarthy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                     Fc-gammaRIIa; human; FCR; model; three-dimension;
                                                                                                                                                                                                                                                                                                                                                                        (first entry
                                                                                                       98US-0099994
98US-0073972
                                                                                                                                                99WO-IB00367
                                                   Garrett TPJ,
                                                                                                                                                                                                                                                       immunoprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 64; 100.0%; Pred. No.
                                       IJ,
                                                                                                                                                                                                                                                                                                                                                                                                                               261 AA
                                     McKenzie IFC, Pietersz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                           Fc-gammaR;
                                                                                                                                                                                                                                                                 inflammatory cell recruitment; RIa; immune function regulation;
                                                                                                                                                                                                                                                      sfc-gammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              6.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 19;
                                                                                                                                                                                                                                                                                           Fc-epsilonR;
                                                Matthews
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                  BR
                                                                                                                                                                                                                                                                                           IgG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                          · IgE;
                                                                                                                                                                                                                                                                                                                   3-D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Three-dimensional structures and models of Fc receptors, useful computer-assisted drug design

in

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q211.78) at position 25, p135 encodes a Ser residu while clone p98/X2 predicts a Thr residue. At position 58, p90

sition 25; p135 encodes a Ser residue Thr residue. At position 58, p90

New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,

etc.

Example 10;

Page

94a; 160pp;

English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT
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Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                             Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIa and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example,
                                                                                                                                                                           N-PSDB; Q21179
                                                                                                                                                                                            WPI; 1992-056864/07
                                                                                                                                                                                                                                                                (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                   13-ЛИІ-1990;
                                                                                                                                                                                                                                                                                                                                      15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                       23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                        W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rapid immunoselection cloning technique; cell surface antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage-specific FcRI receptor encoded by clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R22549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   an FC receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of the FCR can be used to design bioactive chemical compounds, e.g. oligonucleotides, peptides, peptidomimetics and small organic molecules, oligonucleotides, peptidomimetics and small organic molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunodiagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 11; Page 316-317; 326pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       by computer-assisted methods of drug design. Therapeutic compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              210 fymgsktlrgrn
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                                                                                                                                                                                                                             Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; ilarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                   90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA;
                                                                                                                                                                                                                                                                                                                                    90WO-US04986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12
                                                                                                                                                                                                                               Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 64; DB 20;
Pred. No. 0.00019;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        model of an Fc receptor (FcR) protein (3-D) structure that substantially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT
R91439
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Best Local
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macrophages.
coded for a v
of the p98 pr
                                  The amino acid sequence (R91439) of human FCRI was detd. from a CDNA clone, p98 (T14718), obtd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. 2 Other CDNA clones, p135 (T14717) and p90 (T14719),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane spanning domain sequence.
                                                                                                                                                                                                      Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R91439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R91439 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FcRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                     N-PSDB; T14718.
                                                                                                                                                                                                                                                                                          WPI; 1996-200279/20
                                                                                                                                                                                                                                                                                                                                   Aruffo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six
                                                                                                                                                             Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                     18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              229
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3-JUL-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5-FEB-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FYMGSKTLRGRN 12
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                 a variants of the FCRI
                                                                                                                                                                                                                                                                                                                                   Seed B;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                    92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                   93US-0139273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "amino acid 58 is Leu in clone p135
translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         344 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.00025;
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                                                                                                                                                                                                                            useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
               inal sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                domain,
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Y96183
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Best Local s
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and p90 products (see also R91438 and W00859). P of FCRI and a receptor ligand will be helpful in potency of antibodies in therapy.
                                                                                                                    /note=
Misc-difference 126
                                                                                                                                                                                                 /note=
Misc-difference 116
                                                                                                                                                                                                                                                             /note=
Misc-difference 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                  amyloidosis; multiple sclerosis; parasitic disease; Fc
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                                                            Misc-difference
                                                                                 Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                              panning; immunodiagnosis; diagnosis; immunotherapy; gene th
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Macrophage; FcRI; cell surface antigen; human; immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human macrophage-specific FcRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        229
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l Similarity 100.
12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                            /note=
- 82
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51
/note=
213
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123
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117
                                                                                  /note=
134
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                     140
                                                                                                       129
                                                                                                                                                                            /note=
                               /note=
                                                    /note=
                                                                       /note-
                                                                                                                /note-
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                              "encoded
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          encoded by
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Pred. No.
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          AAT"
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. 0.00025;
ches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A fusion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 344;
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229

fymgsktlrgrn 240

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13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                        surfaces other than macrophages, protein of FcRI and a receptor 1
                                                                                                                                                     based upon transient expression of a CSA in eukaryotic
                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                             from a cDNA library expressed in COS cells using a nother invention designed to isolate CSA nucleic acids.
                                                                                                                                                                                                                Example 10; Column
                                                                                                                                                                                                                                   diseases
                                                                                                                                                                                                                                                                                 N-PSDB; A50631
                                                                                                                                                                                                                                                                                                              Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                               29-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                US6111093-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
Misc-difference 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=
Misc-difference 216
                                                                                                                                                                                                                                                                                                                                (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                             28-OCT-1998;
                                                                                                                                                                                                                                          nfections or disorders, e.g. asthma, immune-complex disease,
                                                                                                                                                                                                                                                                                            2000-586382/55.
                                                                                                                                                                                                                                                                                                                                                            89US-0379076
90US-0498809
                                                                                                                                                                                                                                                                                                                                                                               92US-0983647
88US-0160416
                                                                                                                                                                                                                                                                                                              Seed
                                                                                                                                                                                                                                                                                                                                                                                                            98US-0181612
                                                                                                                                                                                                                                                                                                                                                     90US-0553759
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note=
338
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268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note=
220
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                                                                                                                                                                                                              53-55; 75pp;
                                                                                                                                                                                                                                                                                                              В;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded
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                                                                                                                                                                                                             English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAA."
                       in therapy.
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The present sequence is that of a human macrophage specific FcRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of IgG, normally located on the cell surfaces of macrophages. The immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disordinals, including humans. These disorders include asthma, (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the physical selection of cells expressing the antigen by adhesion to the potencies of antibodies ability to interfere with such bonding, or to cause it to occur on proteins they encode, are useful for immunodiagnostic and tor ligand will in therapy. and disorders in The method is to increase

Sequence 344 AA;

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Query Match
Best Local Similarity
----hes 12; Conserv
1 FYMGSKTLRGRN 12
                          Conservative
                                 100.0%;
                       Score 64; DB 21;
Pred. No. 0.00025;
                                         Length 344;
                        Indels
                        0,
                       Gaps
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0;

21-MAY-1992

(first entry)

0

RESULT R22550 ID R2 XX AC R2 XX DT 21 XX В Š RESULT R20811 Query Match Best Local S Matches 12 typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the R22550; R22550 standard; Protein; Sequence This amino acid sequence was predicted from the cDNA sequence of CDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clone predict a Thr residue. At position 58, p135 predicts Leu and ps predicts Val. Sequences predicted from all 3 clones show the predicts Val. Sequences predicted from all 3 clones show the potency New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic Seed B, 15-JUL-1990; 23-JAN-1992 WO9201049-A Homo sapiens Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor. 21-MAY-1992 R20811 standard; Protein; Example 10; Page 94a; 160pp; English 13-JUL-1990; Human (GEHO-) 229 1 FYMGSKTERGRN 12 fymgsktlrgrn 1992-056864/07 macrophage-specific FcRI receptor encoded by clone p135. l Similarity 100 12; Conservative of antibodies in therapy. GEN HOSPITAL CORP Aruffo A, Amiot M; 374 AA; (first entry) 9008-0553759. 90WO-US04986. 240 100.0%; 's 100.0%; 's tive 0; 374 AA 374 Score 64; DB 13; Pred. No. 0.00028; ); Mismatches 0; A it neoplasms, Length 374; Indels for etc. clones of Gaps

parasitic

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Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This amino acid sequence was predicted from the cDNA sequence of cDNA clone p90: It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                           Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                 30-0CT-1996 (first entry)
                                                                                                                                                                                                                                                                        R91438
   Misc-difference
                                                                                                          Homo sapiens
                                                                                                                                                                                               Human FcRI (cDNA clone p135 product).
                                                                                                                                                                                                                                                                                                         R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1992-056864/07
N-PSDB; Q21180.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seed B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Page 94a; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-JAN-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rapid immunoselection cloning technique; cell surface antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human macrophage-specific FcRI receptor encoded by clone p90
                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                 229
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                                                                                                                                                                                                                                                                                                                                                                                                                                  1 FYMGSKTLRGRN 12
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                                                                                                                                                                                                                                                                                                                                                                                             fymgsktlrgrn 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90US-0553759
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                                                    Location/Qualifiers
                /note= "amino acid 25 is Thr in clone p90 and p98
    translated products"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Sc
100.0%; Pr
ntive 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 64; DB 13;
Pred. No. 0.00028;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                                                                                                          immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     etc
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W00859
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Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, pol (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, pol (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The amino acid sequence (R91438) of human FcRI was detd. from a cDNA clone, pl35 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FcRI is a high affinity receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cloning of cDNA encoding cell surface antigen - useful for isolation
                  01-DEC-1992;
25-FEB-1988;
                                                                                                                                       US5506126-A
                                                                                                                                                                                                              Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                           W00859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence. A fusion protein of real and a four in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-200279/20
N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aruffo A,
                                                                      25-FEB-1988;
                                                                                                        09-APR-1996
                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                  Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                     30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                          W00859 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                       (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                      88US-0160416
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93US-0139273
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89US-0379076
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 64; DB 17;
Pred. No. 0.00028;
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Length 374; Indels

0,

Gaps

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RESULT 10
W80448
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Best Local Similarity
Matches 12; Conserv
Misc-difference
                      Misc-difference
                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The amino acid sequence (W00859) of human FCRI was detd. from a cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy
                                              Misc-difference
                                                                                             Misc-difference
                                                                                                                    Misc-difference
                                                                                                                                            Misc-difference
                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                   Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                       W80448;
                                                                                                                                                                                                                                                                                                                                                                                                             W80448 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cloning of cDNA encoding cell surface antigen
                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                              Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                         Human Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-PSDB; T14719.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-200279/20
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18-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       229 fymgsktlrgrn 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                    /note=
77
                                                                                                                                                                                                                     /note=
23
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93US-0139273
                      /note=
141
                                             /note-
103
                                                                      /note=
99
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85
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                                                                                                                                                                                                      "encoded by
                                                                                                                                                                                                                              "encoded by TGG"
                                                                                                                                                      "encoded by CTG"
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                                                                                                                                                                                                                                                                                                                                                                                                            374 AA.:
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0
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Pred. No. 0.00028
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                                                                               by
                                                                                                      bу
        by GGC"
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                                  CCC.
                                                        CAA"
                                                                               TCC"
                                                                                                        AAT"
                                                                                                                                CCC"
                                                                                                                                                                                                       CTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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4444444
                                                                                                                   expression libraries is based on transient expression of an antigen by an in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
Sequence
                                                                                     purified genes and proteins are useful
                                                                                                                                                                                                                                                                                                   from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see X07372) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                                                                                                                                                                                                                                                                                                                            cells.
                                                                                                                                                                                                                                                                                                                                                                                          This is the amino acid sequence of human Fc receptor I (FcRI), deduced from cDNA clone pl35 (see V63456) isolated using a rap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US5830731-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
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                                                                                                        the generation of very large mammalian expression libraries.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-1997;
                                                                      mmunotherapeutic applications,
                                                                                                                                                                                                                                                                                                                                                                      immunoselection cloning method from a cDNA library expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              numan lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1998-609251/51
                                                                                                                                                                                                                                                                                                                                                     The cDNA library was constructed from polyA RNA of cells
                                   including
                                                       0f
                                                    immune-mediated infections,
 A,
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88US-0160416
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176
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256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note=
                                   humans.
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                                                  useful for immunodiagnostic and including the diagnosis and fections, diseases, and disorder
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                                                    disorders
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RESULT
W97833
ID W97
XX
AC W97
                                                                              Query Match
Best Local Similarity
--+-hes 12; Conserv
                                                         Вb
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                 W97833 standard; Protein; 374 AA.
                                                       1 FYMGSKTLRGRN 12
                                                                                        Conservative
                                                                                             100.0%;
                                                                                        0;
                                                                                              Score 64;
Pred. No.
                                                                                        Mismatches
                                                                                              0.00028;
                                                                                                    Length 374;
                                                                                       Indels
                                                                                       0;
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DB 19;

Gaps

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of

W97833;

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New cloning vector and polylinker - based on existing efficient cloning and expression of mammalian cDNA(s), human lymphocyte antigenic sequences
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Misc-difference 159
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Misc-difference 99
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N-PSDB; X07372.
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                                                           cells.
                                                                                                                                  Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                                    Aruffo A,
                                                                                                                                                                                                                                                                                                                            21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1997;
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   expression libraries
                                                        Phis is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p90 (see x07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                (GEHO ) GEN HOSPITAL CORP
                              from a single patient undergoing extracorporeal induction therapy. Clones p135 (see V63456) and
                                                                                                                                                                                                                                                                                                                                                                       25-FEB-1988
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85
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90US-0553759
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176
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23
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                method for cloning cDNAs
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 transient expression
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                                              interleukin-2
                              (see X07373)
of an antigen
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                  from mammalian
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RESULT
W97834
ID W9
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Best Local S
Matches 12
US5830731-A
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                                                                                                                                    Misc-difference
                                                                                                                                                                                           Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   adhesion to
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fymgsktlrgrn
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77
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Pred. No. 0.00028;
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d is useful for
e expressed and
otic cell. It
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                       adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                      Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    induction therapy. Clones pl35 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen by adhesion to an antibody-coated substrate. The method is useful for and an antibody-coated substrate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in Cocells. The cDNA library was constructed from polyA RNA of cells
                                                                          Human macrophage-specific FcRI
                                                                                                   19-DEC-2000
                                                                                                                             Y96134;
                                                                                                                                                     Y96134 standard;
                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                         animals,
                                                                                                                                                                                                                                                                                                                                                                     immunotherapeutic applications, treatment of immune-mediated in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New cloning vector and polylinker - based on existing sequences fefficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aruffo A,
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                                                                                                                                                                                                                                          FYMGSKTLRGRN 12
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DB; X07373.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                       including humans
                                                                                                                                                                                                                                                                                                                                  374 AA;
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                                                                                                                                                                                                                                                                    Conservative
           multiple sclerosis;
                                                                                                (first entry)
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97US-0861205
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89US-0379076
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                                                                                                                                                     Protein;
                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                     374
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                                                                                                                                                                                                                                                                                                                                                                     infections,
                                                                                                                                                                                                                                                                             Score 64; DB 19;
Pred. No. 0.00028;
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                                                                                                                                                                                                                                                                                                                                                                                 including the diagnosis
          parasitic disease; Fc receptor
                                                                                                                                                                                                                                                                                         Length 374;
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                                                                                                                                                                                                                                                                                                                                                                     disorders of
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13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                     Stamenkovic
                                           (GEHO ) GEN HOSPITAL CORP
                                                                                                                        01-DEC-1992;
                                                                                                                                                 28-OCT-1998;
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                                                                    92US-0983647.

88US-0160416.

89US-0379076.

90US-0498809.

90US-0553759.
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216
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306
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305
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139
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126
                     Seed
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213
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134
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136
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116
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2000-586382/55

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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is that of a human macrophage specific PCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in uclaryotic cells and physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                                                                               Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil, HIV; IGG; immunosuppressive; antirheumatic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-SEP-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 10; Column 53-55; 75pp; English.
N-PSDB; A27466
                   WPI; 2000-367968/32
                                                                                                      03-DEC-1998;
                                                                                                                                             03-DEC-1998;
                                                                                                                                                                                      07-JUN-2000
                                                                                                                                                                                                                             EP1006183-A1
                                                                                                                                                                                                                                                                                                             systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                         Human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Y96226;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Y96226 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ability to interfere with such bonding, or surfaces other than macrophages, is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sclerosis.
                                                          (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the potencies of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein of FcRI and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nimals, including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      229 fymgsktlrgrn 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immune-complex disease, amyloidosis, parasitic diseases or multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 FYMGSKTLRGRN 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   rosis. FCRI is a high affinity receptor for the Fc portion of normally located on the cell surfaces of macrophages. The ity to interfere with such bonding, or to cause it to occur on aces other than macrophages, is useful in therapy. A fusion each of FCRI and a receptor ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                               erythematosus;
                                                                                                   98EP-0122969
                                                                                                                                             98EP-0122969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                          FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 These disorders include asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 64;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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glycosylation, cancer
                                                                                                                              receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant portions may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel Fc. receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
                                                                                                                                                                                                                                                                                                                                                                        role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds cells expressing Fc receptors. FcgammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on ff.
                                                  prevent autoimmune diseases, allergies or tumours, especially AIDS rheumatoid arthritis or {\tt MM}.
                                                                                                     compositions containing recombinant proteins may be used to treat or
                                                                                                                                                                                                                                                                                               may be modified to produce recombinant versions. The recombinant Fc
                                                                                                                                                                                                                                                                                                                           neutrophils and eosinophils. Upon Fc receptor activation, immune effecton pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                             FoganmaRI is also known as CD64. Fc receptors play an important role in defending the body against infections. First, pathogens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is the human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 26-28; 60pp; English
                                                                                                                                                                                                                                                                                                                                                          immune effector
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Matches Query Match Best Local ( Human cancer associated protein sequence SEQ ID NO:1128. 08-FEB-2001 В43683; B43683 standard; Protein; 399 AA 15 , 1 FYMGSKTLRGRN 12 l Similarity 12; Conserv Conservative (first entry) 100.0%; 0; Score 64; Pred. No. 0.00028; DB 21; Length 374; Indels 0; Gaps 0

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Sequence

374 AA;

vasotropic; antipsoriatic; antianglogenic; gene therapy; inflammation; immune disorder; haematopoletic cell disorder; autoimmune disorder; allergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; diagnosis; Human; neurological disease; dermatological; naemostatic; cancer associated gene; cancer antigen; detection; cytostatic; neuroprotective; thrombolytic; drug screening proliferative; vulnerary; coagulant; nootropic; immunomodulator; cancer;

Homo sapiens.

XX SO

WO200055350-A1

PR XX

08-MAR-2000; 2000WO-US05882.

(HUMA-) HUMAN GENOME SCI INC

12-MAR-1999;

99US-0124270.

Rosen CA, Ruben SM

WPI; 2000-587533/55

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N-PSDB; C77892
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Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer  $\cdot$ 

Claim 11; Page 1739-1740; 2352pp; English.

cc antityroid; antipleuratic, antipartities, autorities, dermatological; cc neuroprotective; cardiant; thrombolytic; coagulant; nootropic; antipscriatic and antiangiogenic. The polynucleotides and cc vasotropic; antipscriatic and antiangiogenic. The polynucleotides and cc polypeptides can be used for preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosing pathological conditions. Polynucleotides, conditions and diagnosing pathological conditions. Polynucleotides, conditions and the present conditions and the present discorders and antagonists from the present cc invention may be used to treat immune discorders by activating or immune clinibiting the proliferation, differentiation or mobilisation of immune cells, to treat discorders of haematopoletic cells, autoimmune discorders, callergic reactions, graft versus host disease and organ rejection, condulate haemostatic or thrombolytic activity, modulate inflammation, cc cancers, cardiovascular discorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and cc antagonists may be also be used in drug screens. C78449 to C78457 and contraction and contraction contractions are present sequences used in the exemplification of the present formantion. C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in. Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;

## Sequence 399 AA;

Query Match
Best Local Similarity
Matches 12; Conserv 100.0%; Score 64; DB 21; nilarity 100.0%; Pred. No. 0.0003; Conservative 0; Mismatches 0; Length 399; Indels 0; Gaps 0;

1 FYMGSKTLRGRN 12

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Search completed: June 4, 2001, 12:13:29 Job time: 202 sec

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scoring table:
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
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64
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                     185757 segs, 19210857 residues
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Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
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365
393
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US-08-332-562A-94
US-08-080-255-7
US-08-465-713-7
PCT-US93-05857-7
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Sequence
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                                                                                                               Appl
Appl
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## ALIGNMENTS

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TOPOLOGY:
US-08-332-562A-94
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                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                            FILING DATE: 27-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54270/119/GRHA
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5309
TELEPAX: (202)672-5399
                             SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
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                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 07/896,457 FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                904136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HULETT, Mark D. FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                                                                                                                                                                                                                     PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BONADONNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SCOTT, Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HOGARTH, Mark
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               linear
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                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
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                            single
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Best Local Similarity

66.7%;

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RESULT 3
US-08-465-713-7
                                                                                                                              Sequence 7, Application US/08465713 Patent No. 6121419
                                                                                                                                                                                                                                                                                           Matches
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                                                                                                               ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (512) 474-7577 NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LAPPY disk

OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Re-bease #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/080
                                TITLE OF INVENTION: TITLE OF INVENTION:
                                                                          APPLICANT: Rowley, Janet I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Parker, David L.
REGISTRATION NUMBER: 32,165
REGISTRATION NUMBER: AR
REFERENCE/DOCKET NUMBER: AR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 320-7200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: Touston
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ATTORNEY/AGENT INFORMATION:
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ORRESPONDENCE ADDRESS:
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COUNTRY: UL.
77210
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8; Conser
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                                                                                            Rowley, Janet D.
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                                                              COMPOSITIONS AND METHODS FOR
                                DETECTING GENE REARRANGEMENTS AND TRANSLOCATIONS
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                                                                                                                                                                                                                                                                                                         Score 37; DB 1; Pred. No. 48;
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RESULT 4
PCT-US93-05857-7
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GENERAL INFORMATION:
APPLICANT: Board of APPLICANT: The Un
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 320-7200
TELEPAX: (512) 474-7577
NFORMATION TO THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF THE PROPERTY OF
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0
FILING DATE: 17/06/92
                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,713
FILING DATE: 06-UUN-1995
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MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER: US/08/080,255
FILING DATE: 17 JUNE 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                        APPLICATION NUMBER: POPILING DATE: 19930617 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 FTPGSQTERGRN 72
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TOPOLOGY: lir
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
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Similarity 66.7%;
8; Conservative
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THE UNIVERSITY OF TEXAS SYSTEM
TO UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE UP TO THE U
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                                                                                                                                                                                                                                                                                                                                                                                           Patentin Release #1.0,
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Arnold, White & Durkee O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   O. Box 4433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single
                                                        07/900,689
                                                                                                                                                                                                                                                                               PCT/US93/05857
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                                                                                                                                                                                                                                                                                                                                                                                           Version #1.25
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Query Match
Best Local Similarity
...thes 8; Conserv.
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; MOLECULE TYPE: DNA (genomic)
PCT-US93-05857-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity ou.
Thes 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   equence 5, Application US/08061376
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SENERAL INFORMATION:
                                                                                                                                                                                         TELEFAX: (619)546-939
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 1400 amino acids
TYPE: amino acid
                                                                                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Rele
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                     TELEPHONE: (619)546-4737
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 444 SOUTH
CITY: Los Angeles
STATE: California
                                                                                                                   TOPOLOGY:
                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                  NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 FTPGSQTERGRN 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 32,165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Parker, David REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 FYMGSKTLRGRN 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E OF INVENTION: CHARACTERIZATION OF A CHROMOSOME 11Q23
E OF INVENTION: TRANSLOCATION BREAKPOINT ASSOCIATED WITH ACUTE LEUKEMIAS
                                                                                                                                                                                                                                                                                                                                                                                                                                               90071
                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Pretty, Schroeder, Brueggemann & Clark 444 South Flower Street, Suite 2000
                                                                                                                                                         3969 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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Selleri, Licia
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                                                                                                                                                                                                       (619)546-9392
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                                                                                                                                                                                                                                                                                                                                                                                                 IBM PC compatible
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66.7%;
                           57.8%;
66.7%;
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Pred. No.
                           Score 37;
Pred. No.
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                          1.6e+02;
                                           DB 4;
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                                       Length 3969;
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                                                                                                                                                                                                                                                  RESULT
                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 70.0
Matches 7; Conservative
                                                                                                                                                                                                  Sequence 7, Application US/08469667 Patent No. 5733748
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ATTORNEY/AGENT INFORMATION:
NAME: Goldman, Michael L.
REGISTRATION NUMBER: 30,
                                                                                                       NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                              129 MGAKSLIGRN 138
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                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Lactococcus lactis subsp. lactis INDIVIDUAL ISOLATE: ORF2
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                                                           STREET:
                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                         3 MGSKTLRGRN 12
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                                                                                          DDRESSEE:
                                                                                                                                                                                      INFORMATION:
07068-1739
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                                           Roseland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             978450
                                                           6 Becker Farm Road
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                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (716) .263-1487
                                                                                                                                                      Rosen, Craig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Godon, Jean-Jacques
                                                                                                                                                                     Yu, Guo-Liang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ehrlich, Stanislav
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               :: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0, Version #1.25
                                                                          Carella, Byrne, Bain, Gilfillan, Cecchi
Stewart & Olstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (716) 263-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (WUT)
                                                                                                                                       Colon Specific Genes and Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthase from Lactococcus and its applications
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70.0%;
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                                                                                                                                                                                                                                                                                                                                                                          Score 36;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                          DB 1;
11;
                                                                                                                                                                                                                                                                                                                                                                                     Length 259;
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OMPUTER READABLE FORM:

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°CT-US95-07289-7
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GENERAL INFORMATION:
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                                                                        TELEFAX: 201-994-1/44
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO:
            MOLECULE TYPE: protein
                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Ferraro, Gregory D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 32:
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Rosen, Craig
TITLE OF INVENTION: Colon Specific Genes and Proteins
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 135 amino acids
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: 06-JUI
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OPERATING SYSTEM: PC-DOS/MS-DOS
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92 FIGAKDLRGKS 102
                                TOPOLOGY:
                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 07068-1739
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REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/469,667-
FILING DATE: 06-JUN-1995
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SOFTWARE: PatentI
                                                                 LENGTH:
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                                           amino acid
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                                                               135 amino acids
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E: Stewart & Olstein
6 Becker Farm Road
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                                                                                                                                      201-994-1700
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45.5%;
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                                                                                                                                                  RESULT 10
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                                                                                                  Sequence 4, Application US/08341843B Patent No. 5872225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
5; Conserv
                          Patent No. 5872225
                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                        Matches
                                                                                       GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (206) 682-6031
TELEX: 3723836 SEEDANDBERRY
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
TITLE OF INVENTION:
                                                         TITLE OF INVENTION:
                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT:
                                                                                                                                                                                            51 FIMGKKTLQAK 61
                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Seattle
STATE: Washing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0 FILING DATE: 21-APR-1995 CLASSIFICATION: 436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
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Kadlecek, Ann T.
Kadlecek, 39,244
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                                                                                                                                                                                                                                                                                                                                                                                        322 amino acids
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A Method for Characterizing the
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the Nucleotide Sequence
Characterized Thereby
                                           Nucleotide Sequence of L1CAM and
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Pred. No.
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                                                                                                                                                                                                                                                      Mismatches
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UMBER OF SEQUENCES: 3

STREET:

Suite 700

1100 Superior Avenue

Minnich & McKee

Fay, Sharpe, Beall, Fagan,

ADDRESSEE:

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Best Local Similarity
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MEDIUM TYPE: Diskett
MEDIUM TYPE: storabl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OLECULE TYPE: amino acids TYPOTHETICAL: irrelevant
                                                                                                                                                                                                                                                                                                                                                      CLONE: synthesis of several clones UBLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EQUENCE CHARACTERISTICS:
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FILING DATE: June 26, 19
FITORNEY/AGENT INFORMATION:
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56 YSGSFTIEGNN 66
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SOFTWARE: ASCII
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
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                        2 YMGSKTLRGRN 12
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UNDIVIDUAL ISOLATE: 8 day old mouse brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                        I-SENSE:
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44114-2518
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1988
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(216) 9
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                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                       Moos, M.
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                                                                                                                                                                                                                                                                                                                          Tacke, R.
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                                                                                                                                                                                                                                                                               Fruh, K.
                                                                                                                                                                                              NATURE
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5872225ember 18, 1994
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                                                                  Score 32;
Pred. No.
                                                                               DB 2;
                                                                               Length 95;
                                                    Indels
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0;

701-703

JS-08-427-497E-9 Sequence NFORMATION FOR SEQ ID NO: TITLE OF INVENTION: TOPOLOGY: li SEQUENCE CHARACTERISTICS: TELECOMMUNICATION INFORMATION CURRENT APPLICATION DATA: TITLE OF INVENTION: PUBLICATION INFORMATION: HYPOTHETICAL: ATTORNEY/AGENT INFORMATION PRIOR APPLICATION DATA: CORRESPONDENCE ADDRESS: COMPUTER READABLE FORM: MMEDIATE SOURCE: APPLICATION NUMBER: US/08/427,497E FILING DATE: APTIL 24, 1995 MEDIUM TYPE: Diskette MEDIUM TYPE: storable STRANDEDNESS: OPERATING SYSTEM: COUNTRY: ADDRESSEE: ORGANISM: REFERENCE/DOCKET NUMBER: APPLICATION NUMBER: 0 FILING DATE: June 26, IGINAL SOURCE: T-SENSE: JBRARY: NDIVIDUAL ISOLATE: 8 day old mouse brain ELEFAX: ELEPHONE: EGISTRATION NUMBER: ASSIFICATION: 19, A.F. 124 INFORMATION: 44114-2518 OF SEQUENCES: 5969124 Cleveland Minnich, Richard amino acid Application US/08427497E Ohio synthesis of Neural adhesion molecule L1 is a member of the immunoglobulin (216)superfamily Suite (216)U.S.A 1100 Superior Avenue Moos, M. Schachner, M. Scherer, H. Teplow, D. Tacke, R. lamda GT 10 and lamda GT11 NATURE linear Compag Prolinea 5100e (216) 861-5582 irrelevant amino acids 980162 Diskette, single 241-1666 Sharpe, Beall, Fagan, Vance A Method for Characterizing the DOS 5.0 Characterized Thereby the Nucleotide Sequence Nucleotide Sequence of L1CAM and with binding domains several clones 07/904,991 3.50 inch, 149-3-1 720 Kb

; DATE: 1988 US-08-427-497E-9

Best Local Similarity Matches 6; Conserv

Conservative ·

50.0%; 54.5%;

Score 32; DB Pred. No. 23; 1; Mismatches

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Indels

Gaps

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56 YSGSFTIEGNN 66

2 YMGSKTLRGRN 12

Query Match

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JS-09-405-035-1
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Best Local Similarity 66.
Matches 6; Conservative
                                                       Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/09405035 Patent No. 6184351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Biere, Anja Leona
APPLICANT: Citron, Martin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 1
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                                                                                                                                    ORGANISM: ADULT HUMAN BRAIN -09-405-035-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: ADULT HUMAN BRAIN-09-405-035-1
                                                                                                                                                                                                                               SOFTWARE: Patentin Ver. 2.0
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EARLIER FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/405,035
CURRENT FILING DATE: 1999-09-24
EARLIER APPLICATION NUMBER: 60/101,862
EARLIER FILING DATE: 1998-09-25
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TYPE: PRT
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                                                                                                                                                                         TYPE: PRT
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PPLICANT: Citron, Martin
                                                                                                                                                                                                                                                                                                                                              ITLE OF INVENTION: ALPHA-SYNUCLEIN SUPER-MUTANTS ACCELERATE ITLE OF INVENTION: ALPHA-SYNUCLEIN AGGREGATION ILE REFERENCE: A-565
                                                                                                                                                                                           ENGTH:
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2 YMGSKTLRG 10
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66.7%;
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66.7%;
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Pred. No. 36;
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Search completed: June Job time: 223 sec

4, 2001, 12:14:30

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LENGTH: 140
TYPE: PRT
ORGANISM: ADULT HUMAN BRAIN
US-09-405-035-4
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; ORGANISM: ADULT HUMAN BRAIN
US-09-405-035-3
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Best Local Similarity
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Best Local Similarity
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Patent No. 6184351
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Patent No. 6184351
                                                                                                                                                                                                                                                                                  FILE REFERENCE: A-565
CURRENT APPLICATION NUMBER: US/09/405,035
CURRENT FILING DATE: 1999-09-24
EARLIER APPLICATION NUMBER: 60/101,862
EARLIER FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 6
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CURRENT FILING DATE: 1999-09-24
EARLIER APPLICATION NUMBER: 60/101,862
EARLIER FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 6
                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: ALPHA-SYNUCLEIN SUPER-MUTANTS ACCELERATE TITLE OF INVENTION: ALPHA-SYNUCLEIN AGGREGATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE:
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APPLICANT: Citron, Martin
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APPLICANT: Citron, Martin
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39 YVGSKTKEG
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47
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Pred. No. 36;
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Pred. No.
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## ALIGNMENTS

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RESULT 2 A39878 FC gamma (IGG) receptor I-A (high affinity) precursor - human FC gamma (IGG) receptor I-A (high affinity) precursor - human N;Alternate names: CD64 C;Species: Homo saplens (man) C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000 C;Accession: A39878; I70304; B41357; S03018; I57525 R;Van de Winkel, J.G.J; Ernst, L.K.; Anderson, C.L.; Chiu, I.M. J. Biol. Chem. 266, 13449-13455, 1991 A;Title: Gene organization of the human high affinity receptor for IgG, FcgammaRI (CD A;Reference number: A39878; MUID:91302383 A;Accession: A39878 A;Molecule type: DNA A;Residues: 1-374 <van></van>	Oy 1 FYMGSKTLRGRN 12	Query Match 100.0%; Score 64; DB 2; Length 344; Best Local Similarity 100.0%; Pred. No. 6.9e-05; Matches 12; Conservative 0; Mismatches '0; Indels 0; Gaps 0;	A; Molecule type: mRNA A; Molecule type: mRNA A; Residues: 1-344 <all2> A; Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Note: the authors translated the codon ACT for residue 25 as Ala C; Superfamily: Fc gamma receptor I; immunoglobulin homology C; Superfamily: alternative splicing; immunoglobulin receptor; transmembrane protein C; Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein F; 117-170/Domain: immunoglobulin homology <imm></imm></all2>	A; Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 A; Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 R; Allen, J.M.; Seed, B. Nucleic Acids Res. 16, 11824, 1988 Nucleic Acids Res. 16, 11824, 1988 A; Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (A; Reference number: S03018; M0ID:89098339 A; Accession: S03019	R;Allen, J.M.; Seed, B. Science 243, 378-381, 1989 A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284 A;Accession: A41357 A;Molecule type: mRNA	RESULT 1 A41357 FC gamma (IgG) receptor I (high affinity) form b - human N.Alternate names: CD64 C. Species: Homo sapiens (man) C. Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000 C. Accession: A41357: S03019

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Science 277,
                                       A;Reference number: A64720; MUID:97426617
A;Accession: H64726
                                                                                                                       R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, A.; Rose, D.J.; Mau, B.; Shao, Y.
                                                                                                                                                                                                                                abP protein -
                                                                                                                                                                   Species: Escherichia coli
,Date: 12-Sep-1997 #sequence_revision 17-Sep-1997 #text_change 04-Mar-2000
,Accession: H64726
                   Status: nucleic acid sequence not shown;
                                                                              Title: The complete genome sequence of Escherichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wolecule type: mRNA; Wolstidues: 1-24, 'S, '26-337, 'T, '39-374 <AL2>; Residues: 1-24, 'S, '26-337, 'T, '39-374 <AL2>; Cross-references: EMBL:X14356; NID:93133; Perez, C.; Wietzerbin, J.; Benech, P.D. ol. Cell. Biol. 13, 2182-2192, 1993 ol. Cell. Biol. 13, 2182-2192, 1993; Title: Two cis-DNA elements involved in mye
                                                                                                                                                                                                                                                                                                                                                                                                                                               Best
Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reference number: Accession: 157525
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Cross-references: GB:X14356; GB:M21091; NID:g31331;
Allen, J.M.; Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GB:L03418; Allen, J.M.; Seed, B.
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Residues: 1-8 <RE2>
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293-313/Domain: transmembrane #status predicte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Keywords: alternative splicing; glycoprotein;
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Reference number: S03018; M
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                                                                                               1453-1462, 1997
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Pred. No. 7.5e-05;
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Best Local
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Best Local
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C; Superfamily: Fc gamma
                                                                                                                                                                                                                                                                                                                                                                                                        N;Alternate names: high affinity IgG re
C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Linkage on chromosome 3 of autoimmune diabetes A;Reference number: 148471; MUID:93242399
A;Accession: 148471
A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology
                                                                                                                      A; Title: Molecular cloning and expression of A; Reference number: A43511; MUID: 90111035
                                                                                                                                                                                                                                                                                              3;Osman, N.; Kozak, .....
J. Immunol. 148; 1570-1575, 1992
A;Title: Structure and mapping of the gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  F;128-180/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:X70980; NID:g311748; PIDN:CAA50311.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Reywords: immunoglobulin receptor
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A; Residues: 1-336 < RES>
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C; Superfamily:
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A; Experimental source
                                                             A; Molecule type: mRNA
                                                                              A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                         C; Accession: A46480;
                                                                                                     A; Accession: A43511
                                                                                                                                                                                                                            \;Molecule type: DNA
\;Residues: 1-404 <0
                                                                                                                                                                                                                                                                    1;Status: preliminary
                                                                                                                                                                                                                                                                                        A; Accession: A46480
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                                                                                                                                                                                  NOTE: Sequence extracted from NCBI backbone Sears, D.W.; Osman, N.; Tate, B.; McKenzie,
                                                                                                                                                                                ;Sears, D.W.; Osman, N.;
                                                                                                                                                                 Immunol. 144,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 FYMGSKTLRGRN
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95~698, 1993
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                                                                                                                                                                 sman, N.; Tate, B.; 371-378, 1990
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                                                                                                                                                                                                                                                                                                                                                                    C.A.; McKenzie, I.F.; Hogarth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GB:AE000116; GB:U00096; NID:g1786240; PIDN:AAC73167.1; PID:g17862
ce: strain K-12, substrain MG1655
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54.5%;
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Pred. No. 2.4;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 40; DB Pred. No. 3.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    receptor
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                                                                                                                                                                                                                                                                                                                                                                                                            18-Nov-1994 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-Jul-1996 #text_change 21-Jan-2000
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2.4;
                                                                                                                                     the mouse high affinity Fc receptor
                                                                                                                                                                                  (NCBIN:85205, I.F.C.; Hogar
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                                                                                                                                                                                                       NCBIN:85208,
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                PID:g200753
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C;Date: 14-Aug-1998 #sequence_revision 14-Aug-1998 #text_change 05-Nov-1999
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                 В
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A;Accession: H71147
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A; Residues: 1-379 <K
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A;Accession: E71162
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DNA Res. 5, 55-76, 1998
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                                                                                                                                                                                                                                                            A;Residues:
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                                                                                                                                                                                                                                                                                        A; Status: preliminary; nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                 C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 hypothetical protein PHO394 - Pyrococcus horikoshii
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                                                                                                                                                                                                                                                                                                                                                            M.; Ohfuku, Y.; Funahashi,
NA Res. 5, 55-76, 1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;Date: 10-Sep-1999
;Accession: E71162
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Best Local
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                                                                                                                                                                                                                          Experimental source:
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Matches
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                 264 FYAGSKVFRG
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-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 21-Ju1-2000
                                                                                                                                                                                                                                                        1-276 <KAW>
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                                                                                   Similarity 7; Conserv
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8; Conser
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                                                                                   Conservative
               273
                                                                                                                                                                                                      replaces an interim accession
                                                                                                                                                                                                                        strain OT3
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66.7%;
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                                                                                                 Score 38;
Pred. No.
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Pred. No. 6.9;
                                                                                                                                                                                                                                                                                      sequence not shown; translation
                                                                                 Mismatches
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4.7;
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                                                               A; Status: nucleic acid sequence not shown; not compared with conceptual translation
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                                 A; Residues:
                                                 A; Molecule type:
A;Cross-references: GB:L04284; NID:g184393; PIDN:AAA58669.1; PID:g18439
A;Note: sequence extracted from NCBI backbone (NCBIP:117729)
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Cell 71, 691-700, 1992
A;Title: Involvement of a homolog of Drosophila trithorax A;Reference number: A44265; MUID:93046667
A;Accession: A44265
                                                                                 C;Species: Homo sapiens (man)
C;Date: 30-Apr-1993 #sequence_revision 18-Nov-1994 #text_change
C;Accession: A44265; A44264; I59112; I37165; I38485
R;Tkachuk, D.C.; Kohler, S.; Cleary, M.L.
                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Thehes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R:Chung, S.O.; Kubo, T.; Natori, S.
J. Biol. Chem. 270, 4624-4631, 1995
A;Title: Molecular cloning and sequencing of arylphorin-binding A;Reference number: A56097; MUID:95181458
A;Accession: A56097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              arylphorin-binding protein precursor - C;Species: Sarcophaga peregrina C;Date: 03-Oct-1995 #sequence_revision
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:AE000274; GB:U00096; NID:g1788089; PIDN:AAC74868.1; PID:g17880
A;Experimental source: strain K-12, substrain MG1655
C;Superfamily: hypothetical protein b1798
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Science 277, 1453-1462, 1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 1-1163 <C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-212 <BLAT>
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                                                                                                                                                                       N;Alternate names: acute lymphoblastic leukemia gene 1 protein (ALL-1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: GB:D29741
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                                                                                                                                                                                             trithorax homolog HTX, version 2 - human
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Best Local
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8; Conserv
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53.3%;
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Pred. No.
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1-3968 <TKA> mRN?

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; Marschalek, R.; Greil, J.; Lochner, K.; Nilson, I.; Siegler, G.; Zweckbronner, Title: Molecular analysis of the chromosomal breakpoint and fusion transcripts; Reference number: 137165; MUID:95315013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11. /1. /Ul-/Vo, 1996
Title: The t(4):11) chromosome translocation of human seference number: A44264; MUID:93046668
Accession: A44264; Accession: A44264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Status: not compared with conceptual translation
                                                                                   leuD 3'-region hypothetical protein - Lactococcus lactis subsp. lactis
C;Species: Lactococcus lactis subsp. lactis
C;Date: 03-Oct-1995 #sequence_revision 03-Oct-1995 #text_change 17-Mar-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross references: GB:L04731; NID:g339921
Note: sequence extracted from NCBI backbone (NCBIP:117779)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ancer Res. 54,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cross-references: EMBL:X83604; NID:g897757; PIDN:CAA58584.1; PID:g899268 Note: submitted to the EMBL/GenBank/DDBJ databases by R. Marschalek, 20 (GU, Y.; Alder, H.; Nakamura, T.; Schichman, S.A.; Prasad, R.; Canaani, O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Molecule
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cross-references:
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Best Local Similarity
Matches 8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                  1569-1596/Region: zinc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene: GDB:MLL; HTX;
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residues: 1212-1603,'GTE'.
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esidues: 1317-1700,'DD',1703-1936,'H',1938-2180,'S',2182-2328 <DJA>
ross-references: GB:L01986; NID:9307522; PIDN:AAA92511.1; PID:9553800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          itle: A trithorax-like gene is interrupted by chromosome 11q23 sference number: I58112; MUID:93265134
               Title: Gene
                                                                   Accession: F36889
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                           Godon, J.J.; Delorme, C.; Bardowski, J.; Chopin, M.C.; Ehrlich, S.D.; Renault, Bacteriol. 175, 4383-4390, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                    527-1556/Region: zinc
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LQIESTSP',2555-3554,'N',3556-3594,'V',3596-3899,'A',3901-3968
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1992
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990, 1993
in Lactococcus
; MUID:93322316
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Pred. No. 2e+0
1; Mismatches
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                                                                                                                               Lactococcus lactis subsp. lactis (strain IL1403)
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                     lactis:
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2e+02;
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                           amino acid biosynthes:
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Best Local
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C;Keywords: ATP; P-loop
F;19-219/Domain: ATP-binding cassette homology <ABC>
F;36-43/Region: nucleotide-binding motif A (P-loop)
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A; Residues: 1-259 <GOD>
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A; Status: preliminary;
A; Map position: 3
A; Introns: 19/1;
A; Note: C0410w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              J. Bacteriol. 174, 6580-6589, 1992
A;Title: Branched-chain amino acid biosynthesis
A;Reference number: S35132; MUID:93015710
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      hypothetical protein (leuD 3' region) - Lactococcus lactis subsp. lactis
C;Species: Lactococcus lactis subsp. lactis
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A; Residues: 1-259 <GOD>
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                                                                                                                                                                                                                        hypothetical protein C0410w - malaria F
C;Species: Plasmodium falciparum
C;Date: 15-Oct-1999 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Keywords: ATP; P-loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Godon, J.J.;
                                                                                                                                                                    R;Lawson, D.; Bowman, S.; Barrell, submitted to the EMBL Data Library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Superfamily: unassigned
                                                                                                                                                                                                          C; Accession: T18443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Species: Lactococcus lactis subsp. la; Species: 11-Dec-1993 #sequence_revision.
                                                                    A; Cross-references:
                                                                                                      A; Molecule type:
                                                                                                                       A;Status:
                                                                                                                                     A; Accession: T18443
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                                                                                      A;Residues:
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Best Local
                                                                                                                                                         Reference number:
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                                                       Genetics:
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                                                                                                                     preliminary; translated from GB/EMBL/DDBJ
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7; Conservative
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                                                                                                          DNA
                   61/3; 87/2; 117/2; 145/3; 171/2
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                                                                        EMBL: 298547; NID: e1325376; PID: e1407850;
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Y, August 1997
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A;Experimental source: cultivar Williams
C;Superimmily: GTP-binding regulatory protein Gs alpha chain
C;Keywords: GTP binding; signal transduction
                                                                                                                                                                                                                                                                                                                 C;Accession: T07159

R;Kim, W.Y.; Cheong, N.E.; Lee, D.C.; Je, D.Y.; Bahk, J.D.; Cho, M.J.; Lee, S.Y.

Plant Physiol. 108, 1315-1316, 1995

A;Title: Cloning and sequencing analysis of a full-length cDNA encoding a G protein alph
A;Reference number: Z15966; MUID:95357428
                                                                                                                                                                                                                                                                                                                                                                                                                                GTP-binding regulatory protein alpha chain 2 - soybean
C.species: Glycine max (soybean)
C.bate: 30-Apr.1999 #sequence_revision 30-Apr.1999 #text_change 11-May-2000
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submitted to the EMBL Data Library, November 1994
A;Description: pUB2380 carries a putative transposable element (IS1294) capable of medic
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Best Local Similarity 58.3

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Cross references: EMBL:x82430; NID:g572684; PID:g572685
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Residues: 1-385 <KIM>
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www=ncp://www.ncpi.nim.nin.gov/piow/cd/cdo4.ncm.	-i- DATABASE: NAME=PROW: NOTE=CD guide CD64 entry:	-1- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.	-I- TISSUE SPECIFICITY: MONOCYTE/MACROPHAGE-SPECICIC.	PRODUCTS: 2	TYPE I MEMB	•	-!- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS GAMMA. HIGH	Science 243:378-381(1989).		"Isolation and expression of functional high-affinity Fc receptor	MEDLINE-89100284; PubMed-2911749;	SEQUENCE FROM N.A.		Nucleic Acids Res. 16:11824-11824(1988).	•	"Nucleotide sequence of three cDNAs for the human high affinity Fc	Allen J.M., Seed B.;	MEDLINE=89098339; PubMed=2974947;	SEQUENCE FROM N.A.			Catarrhini; Hominidae;	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Homo sapiens (Human).			UL-UCI-2000 (REI. 40, LASC AMMOCALION UPUACE)	(Rel. 12,	(Rel. 12,	315;	FCG1_HUMAN STANDARD; PRT; 374 AA.	_HUMAN	

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CC or send an email to license@isb-sib.ch).
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CC rembl: X14355; CAA32537.1; -.
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P39220; P75632;

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Best Local
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      Science [3]
                                Blattner F.R., Plunkett G. III, Bloch C.A., Riley M., Collado-Vides J., Glasner J.D., Rc Gregor J., Davis N.W., Kirkpatrick H.A., Goe Mau B., Shao Y.;
                                                                   SEQUENCE FROM N.A.
STRAIN=K12 / MG1655;
MEDLINE=97426617; PubMed=9278503;
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              "The complete genome sequence of Escherichia coli K-12.", Science 277:1453-1474(1997).
                                                                                                                   Isono K., Mizobuchi K., Nakata A., Fujika N., "Systematic sequencing of the Escherichia coli genome: analysis the 0-2.4 min region.";
                                                                                                                                               MEDLINE=92334977; PubMed=1630901;
Yura T., Mori H., Nagai H., Nagat
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                                                                                                                                                                                                                                                                                                                                    FYMGSKTLRGRN 240
                                                                                                                                                                                                                                                                                                                                                                          l Similarity
12; Conserv
                                                                                                            Acids
                                                                                                                                                                                                                                                                                                                                                                                                                       374 AA;
                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; Receptor; Transmembrane; Glycoprotein; Signal;
                                                                                                      region.";
Res. 20:3305-3308(1992).
                                                                                                                                     Nagai H., Nagata T.,
hi K., Nakata A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       292
313
374
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Last annotation update)
PROTEIN IN DJLA-RLUA INTERGENIC
                                                                                                                                                                                                              gamma subdivision;
                                                                                                                                                                                                                                                                                                                                                                                                                       MW;
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
BY SIMILARITY.
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S -> T (IN CAA32536).
2C2AA8103ECF16E6 CRC64;
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CYTOPLASMIC (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                  Score 64;
Pred. No.
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                                       H.A., Goeden M.A.,
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                                                                                                                                                                                                                                                                                        216 AA
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                                                 Rode C.K., Mayhew
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(POTENTIAL).
(POTENTIAL).
(POTENTIAL).
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ayhew G.F.,
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Best Local
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MEDLINE-90111035; PubMed-
Sears D.W., Osman N., Ta
"Molecular cloning and e
"Molecular cloning and e
                                                                        modified and this statement is not ren
entities requires a license agreement
or send an email to license@isb-sib.ch
                                                                                                                                                                                                                                                                                                                                                                                                                   OSMAN N., KOZAK C.A., MCKENZIE I.F., HOGATH P.M.;
"Structure and mapping of the gene encoding mouse high affinity
gamma RI and chromosomal location of the human Fc gamma RI gene
J. Immunol. 148:1570-1575(1992).
-!- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS GAMMA. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mus musculus (Mouse).
Mus musculus (Mouse).
'``rvota; Metazoa; Chordata; (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FCG1_MOUSE P26151;
                                                                                                                                                                               the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                      -i- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-i- TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
-i- SIMILARITY: CONTAINS, 3 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
-i- CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-10 IS THE INITIATOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; D10483; -; NOT_ANNOTATED_CDS.
EMBL; AE000116; AAC73167.1; -.
ECOGene; EG12610; Yabp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIGH AFFINITY IMMUNOGLOBULIN GAMMA RI) (FCRI) (IGG FC RECEPTOR I).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=92166399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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15-JUL-1999
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                                                                                                                                                        non-profit
                                                                                                                                                                                                           the Swiss Institute of Bioinformatics and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Rel. 22, Last sequence update)
(Rel. 38, Last annotation updat
IY IMMUNOGLOBULIN GAMMA FC RECER
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                                                                                                                               institutions as long atement is not removed.
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54.5%;
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Pred. No.
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                                                                                                                                                                                                                                    It is produced through a collaboration
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RECEPTOR
                                                                                                    (See http://www.isb-sib.ch/announce/
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                                                                                                                                                ormatics and the EMBL outstation There are no restrictions on it ing as its content is in no wa
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e mouse
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high affinity
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EMBL; M31314; AAA40056.1;

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Best Local Similarity
Matches 8; Conser
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DOMAIN
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SEQUENCE
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DOMAIN
                   InterPro;
InterPro;
                                                                                                                                                                                                                                                                                                                       01-OCT-1996 (Rel. 34, Last annot 60S ACIDIC RIBOSOMAL PROTEIN PO. LIPO-A AND LIPO-B.
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                                      EMBL; X72714; CAA51264.1; -. EMBL; X72714; CAA51263.1; -.
                                                                               modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                 OF E.COLI PROTEIN L10.

-i- SUBUNIT: PO FORMS A PENTAMERIC COMPLEX BY INTERACTION WITH DIMERS OF P1 AND P2.

-i- SIMILARITY: BELONGS TO THE L10P FAMILY OF RIBOSOMAL PROTEINS.
                                                                                                                                                                                                                                                          STRAIN-LEM 75
                                                                                                                                                                                                                                                                                      Leishmania infantum.
Eukaryota; Euglenozoa;
NCBI_TaxID=5671;
                                                                                                                                                                                                                                                                                                                                                               RLAO_LEIIN STA
P39097;
01-FEB-1995 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DOMAIN
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                                                                     or send an email to license@isb-sib.ch).
                                                                                                             the European Bioinformatics Institute.
                                                                                                                                                                                            "Isolation, characterization and analysis of the Leishmania ribosomal PO protein genes."; MO1. Biochem. Parasitol. 61:265-274(193).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00047; ig;
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MGD; MGI:95498; Fcgr1.
                                                                                                                                                                                                                                       Soto M.,
                                                                                                                                                                                                                                                 MEDLINE=94088674; PubMed=8264730;
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IgG-binding protein; Receptor;
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                                                                                                non-profit institutions as long
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Ribosomal_L10;
        60s_ribosomal;

    Last sequence update)
    Last annotation updat

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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
N-LINKED (GLCNAC.) (
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Pred. No.
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(GLCNAC...)
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(POTENTIAL)
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                                                                              .isb-sib.ch/announce/
                                                                                                             restrictions on
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RESULT 6
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use by non-profit institutions as long a
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or send an email to license@isb-sib.ch).
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Eukaryota; Metazoa; Nemat
Rhabditidae; Peloderinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ribosomal protein; Phosphorylation SEQUENCE 323 AA; 34771 MW; 1496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WormPep; C05D10.4; CE01127.
Hypothetical protein.
SEQUENCE 371 AA; 41906 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
HYPOTHETICAL 41.9 KDA PROTEIN COSD10.4 IN CHROMOSOME
                SEQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-97426617; PubMed-9278503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C05D10.4.
Blattner F.R., Plunkett G.
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STRAIN-BRISTOL N2;
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                                                                                                                                                            Proteobacteria; gamma subdivision; Enterobacteriaceae;
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Pred. No.
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Mismatches
Bloch C.A.,
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MBL outstation -
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SEQUENCE
                      viral early region 4 gene.";
Breast Cancer Res. Treat. 39:57-67(1996)
                                                                                 Javier R.,
                                                                                                                                                                NCBI_TaxID=10527;
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EARLY E4 34
      -!- SIMILARITY:
                                                                                                    MEDLINE=96327849; PubMed=8738606
                                                                                                                         SEQUENCE FROM N.A.
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15-DEC-1998
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                                                          "Mammary tumors induced by human adenovirus
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r J., Davis N.W., Kirkpatrick
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Mol. Genet. 7:1703-1712(1998).
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MEDLINE=94215165; PubMed=6162575;
MEDLINE=94215165; PubMed=6162575;
Gu Y. Alder H., Nakamura T., Schichman S.A., Präsad
Saito.H., Croce C.M., Canaani E.;
"Sequence analysis of the breakpoint cluster region
involved in acute leukemia.";
Cancer Res. 54:2326-2330(1994).
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                                                                                                                                                       "The human MIL gene: nucleotide sequence, homology to the Drosophila trx zinc-finger domain, and alternative splicing."; DNA Cell Biol. 14:475-483(1995).
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MEDLINE-93046607; PubMed-1423624;
MEDLINE-93046607; PubMed-1423624;
Tkachuk D.C., Kohler S., Cleary M.L.;
"Involvement of a homolog of Drosophila trithorax by 11q23
chromosomal translocations in acute leukemias.";
                                                                   MEDLINE=95315013; PubMed=7794749;
Marschalek R., Greil J., Lochner K., Nilson
                                                                                                                                                                                                                                                                       SEQUENCE_OF 1251-1654 FROM N.A. (ISOFORM 14P-18B)
MEDLINE-95322025; PubMed-7598802;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Exon/intron structure of the human ALL-1 (MLL) gene involved in translocations to chromosomal region 11q23 and acute leukaemias."; Br. J. Haematol. 93:966-972(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human)
Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MLL OR HRX OR ALL1 OR TRX1 OR HTRX.
                                                                                                                                                                                                                             Rowley J.D.,
                                                                                                                                                                                                                                                    Mbangkollo D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=93265134; PubMed=1303259;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A. MEDLINE=96290553; PubMed=8703835;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                    Neckbronner I., Beck J.D., Fey G.H.;
Molecular analysis of the chromosomal breakpoint and fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE OF 1317-2328 FROM N.A.
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                                                                                                                  EQUENCE OF 1212-1603 FROM N.A.
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Odera Y., Nakazawa S., Ariyoshi Y., Takahashi T., Ueda R.;
Two distinct portions of LTG19/ENL at 19p13 are involved in t(11;19)
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by chromosome 11q23
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SEM cell line with chromosomal
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-i- DISEASE: INVOLVED IN ACUTE LEUKEMIAS BY CHROMOSOMAL TRANSLOCATIONS
T(11;19)(Q23;P13.3) THAT INVOLVES MLL AND MLLT1/ENL;
T(4;11)(Q21;Q23) THAT INVOLVES MLL AND MLLT2/AF4; T(9;11)(P22;Q23)
THAT INVOLVES MLL AND MLLT3/AF9; T(6;11)(Q27;Q23) THAT INVOLVES
MLL AND MLLT4/AF6; T(11;17)(Q23;Q21) THAT INVOLVES MLL AND
MLLT6/AF17; T(x;11)(Q13;Q23) THAT INVOLVES MLL AND MLLT7/AFX1;
T(10;11)(P12;Q23) THAT INVOLVES MLL AND MLLT10/AF10;
T(1;11)(Q21;Q23) THAT INVOLVES MLL AND AF1Q; T(11;19)(Q23;P13.3)
THAT INVOLVES MLL AND ELL; AND T(11;19)(Q23;P23) THAT INVOLVES MLL
AND AF1Q; T(11;19)(Q23;P13.3)
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Proc. Natl. Acad. Sci. U.S.A. 97:2814-2819(2000).
-i- FUNCTION: POSSIBLY ACTS AS A TRANSCRIPTIONAL REG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CHROMOSOMAL TRANSLOCATION WITH GAS7.
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"A method for identifying genes within yeast artificial chromosom application to isolation of MLL fusion cDNAs from acute leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE OF 1421-1540 FROM N.A. MEDLINE-94020842; PubMed-8414518;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dncogene 8:3157-3160(1993)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the Environment of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institute of the Swiss Institut
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VRIIPSSKRTDATIAKQLLQRA -> VSSLILNWKSPRKSG
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A.T HOOK (BY STM)
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RESULT

FUNCTION: GUANINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE

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AC QO2151
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01-FEB-1996 (Rel. 3
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GBA1_SOYBN
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Q02151;
Q1-JUL-1993
Q1-JUL-1993
Q1-JUL-1993
Q1-JUL-1998
"Cloning and sequencing analysis of a full-length protein alpha subunit, SGAI, from soybean."; Plant Physiol. 108:1315-1316(1955).
                                                                                                                                                                                                    GUANINE NUCLEOTIDE-BINDING GPA1 OR GA1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00005; ABC_tran; 1.

PROSITE; PS00211; ABC_TRANSPORTER; 1.

Hypothetical protein; ATP-binding; Transport.

NP_BIND 36 43 ATP (BY SIMILARITY).

NP_BIND 36 43 ATP (BY SIMILARITY).

SEQUENCE 259 AA; 29713 MW; 49EED150F6A15169 CRC64;
                                                                 Kim W.Y.,
                                                                             STRAIN=CV. WILLIAMS;
MEDLINE=95357428; Pu
                                                                                                         SEQUENCE FROM N.A
                                                                                                                                  NCBI_TaxID=3847;
                                                                                                                                               Magnoliophyta; eudicotyledons; core eudicots; Rosidae; Fabales; Fabaceae; Papilionoideae; Glycine.
                                                                                                                                                                        Eukaryota; Viridiplantae;
                                                                                                                                                                                       Glycine max (Soybean).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Godon J.-J., Chopin M.-C., Ehrlich S.D.; "Branched-chain amino acid biosynthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JUL-1993 (Rel. 26, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
15-DEC-1998 (Rel. 37, East annotation update)
HYPOTHETICAL ABC TRANSPORTER ATP-BINDING PROTEIN IN LEUD 3'REGION.
Lactococcus lactis (subsp. lactis) (Streptococcus lactis).
Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              subsp. lactis."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=NCDO 2118;
MEDLINE=93015710; PubMed=1400210;
                                                   Lee S.Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; U92974; AAB81917.1; -.
                                                                                                                                                                                                                                                                                                                                                      129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=1360;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                   3 MGSKTLRGRN 12
||:|:| |||
29 MGAKSLIGRN 138
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        S35136;
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                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 7; Conser
                                                               Cheong N.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IPR001617; -
                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                   STANDARD;
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40,
                                                             PubMed=7630957;
                                                                                                                                                                                                                                                                                                                                                                                                                     56.2%;
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Last annotation update)
WDING PROTEIN ALPHA-1 SUBUNIT (GP-ALPHA-1).
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                                                                                                                                                                        Embryophyta; Tracheophyta;
                                                                                                                                                                                                                                                                                                                                                                                                       N
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 36;
Pred. No.
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                                                                                                                                                                                                                                                                                    PRT;
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                                                                Je D.Y.,
                                                                                                                                                                                                                                                                                385 AA.
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                                                                 Bahk J.D.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                Length 259
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                                       cDNA encoding
                                                                                                                                                           Spermatophyta;
                                                                Cho M.J.,
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Best Local S
Matches 7
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                                                                                                                     Hayward D.C., Delaney S.J., Campbell H.D., Ghysen A., Benzer S., Kasprzak A.B., Cotsell J.N., Young I.G., Miklos G.L.G.; "The Sluggish-A gene of Drosophila melanogaster is expressed in the nervous system and encodes proline oxidase, a mitochondrial enzyme involved in glutamate biosynthesis."; Proc. Natl. Acad. Sci. U.S.A. 90:2979-2983(1993).
                                                                                                                                                                                                                                                                                                                                          01-OCT-1993 (Rel. 27, Created)
01-OCT-1993 (Rel. 27, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
PROLINE OXIDASE, MITCCHONDRIAL PRECURSOR (EC 1.5.3.-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commen
                                                                                                                                                                                                                                                                                                                                                                                Q04499; 061349;
01-0CT-1993 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                     MEDIINE-98188272; PubMed-9520435;
Maleszka R., de Couet H.G., Miklos G.L.G.;
"Data transferability from model organisms to humufrom functional genomics of the flightless region Proc. Natl. Acad. Sci. U.S.A. 95:3731-3736(1998).
                                                                                                                                                                                                                  STRAIN-CANTON-S;
                                                                                                                                                                                                                                                                              Pterygota; Neoptera; Endopterygota; Diptera;
                                                                                                                                                                                                                                                                                                                      SLGA OR SLG.
                                                                                                                                                                                                                                                                                                                              DEHYDROGENASE) (PROTEIN SLUGGISH-A)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                      MEDLINE=93219408; PubMed=8096642;
                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                    Ephydroidea;
                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Arthropoda; Tracheata;
                                                                                                                                                                                                                                                                                                        Drosophila melanogaster (Fruit fly)
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InterPro; IPR002976;
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SUBUNIT: G PROTEINS ARE C
THE ALPHA CHAIN CONTAINS
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                                                                                                                                                                                                                                                                   Drosophilidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Transducer; ADP-ribosylation
           CONVERTS PROLINE TO DELTA-1-PYRROI
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                                                                                                                                                                                                                                                                                                                                                                                                            STANDARD;
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58.3%;
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ADP-RIBOSYL[1] (BY ACTION O
W; 655478036330FFAA CRC64;
                                                                                                                                                                                                                                                                 Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 36;
Pred. No.
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                                                                                                                                                                                                                                                                              Hexapoda; Insecta;
a; Brachycera; Muscomorpha;
                                 region of
                                   to human beings: insights region of Drosophila.";
ROLINE-5-CARBOXYLATE. SLUGGISH BEHAVIOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 385
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                                   This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EW the European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL;
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EMBL; AF044218;
                       or send an
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Query Match
Best Local Similarity
Matches 6; Conser
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042783;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
NEUTRAL TREHALASE (EC 3.2.1.28) (ALPHA, ALPHA-TREHALASE)
                                                                                                               Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databases -:- FUNCTION: POSSIBLY INVOLVED IN MOBILIZATION OF CYTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; L07330; AAA02748.1; EMBL; AF017777; AAC28410.
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                                                                                                                                                                                SEQUENCE FROM N.A.
STRAIN-74-OR23-1A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF01619; Pro_dh; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                                                                D'Enfert C., Bonini
                                                                                                                                                                                                                                                   Sordariales;
                                                                                                                                                                                                                                                                   Eukaryota;
                                                                                                                                                                                                                                                                                     Neurospora crassa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CONFLICT
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                                                                                                                                               erenzi H.F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  interPro;
                                                                                                                                                                                                                                                                                                                       ALPHA, ALPHA-TREHALOSE GLUCOHYDROLASE)
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TISSUE SPECIFICITY: MOST ABUNDANT IN DEVELOPING NERVOUS
PREVELOPMENTAL STAGE: EXPRESSED IN DEVELOPING EMBRYO AS I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     by non-profit institutions as long
                                 SIMILARITY: BELONGS
                                                            CATALYTIC ACTIVITY: ALPHA, ALPHA-TREHALOSE + H(2)0 = 2 D-GLUCOSE. SUBCELLULAR LOCATION: CYTOPLASMIC (BY SIMILARITY).
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                                            TO FAMILY 37 OF GLYCOSYL HYDROLASES
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                                                                                                                                                                Zapella P.D.A.,
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                                                                                                                                                                                                                                                   Neurospora
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 36;
Pred. No.
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collaboration
L outstation -
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email to license@isb-sib.ch)

(See http://www.isb-sib

and

commercia

EMBL outstation

InterPro; IPR001661; -. Pfam; PF01204; Trehalase;

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                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Exthe European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                                                            Q59189;
15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
TOPOISOMERASE IV SUBUNIT B (EC.5.99.1.-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peterson J., Kerlavage A.R., Quackenbush J., Salzberg S., Hanson M., van Vugt R., Palmer N., Adams M.D., Gocayne J.D., Weidman J., Utterback T., Watthey L., McDonald L., Artiach P., Bowman C., Garland S., Fujii C., Cotton M.D., Horst K., Roberts K., Hatch B., Smith H.O., Vootar T.
                                                                                                                                                                                                                                                                                                   Ojaimi C., Davidson B.E., Saint-Girons I., Old I.G., "Conservation of gene arrangement and an unusual organization genes in the linear chromosomes of the Lyme disease spirochaet Borrelia burgdorferi, B. garnini and B. afzelli.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fraser C.M., Casjens S., Huang W.M., Sutton G.G., Clayton Lathigra R., White O., Ketchum K.A., Dodson R., Hickey E.F. Dougherty B., Tomb J.-F., Fleischmann R.D., Richardson D., Dougherty B., Tomb J.-F., Fleischmann R.D., Richardson D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BORBU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS; PR00744; GLHYDRLASE37
                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=95111614; PubMed=7812434;
                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE OF 1-83 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                Nature 390:580-586(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                burgdorferi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Smith H.O., Venter J.C.; "Genomic sequence of a Lyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=98065943; PubMed=9403685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Borrelia burgdorferi (Lyme disease spirochete)
Bacteria; Spirochaetales; Spirochaetaceae; Bor
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SEQUENCE 728 AA; 84121 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS00927; TREHALASE_1; PROSITE; PS00928; TREHALASE_2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               271
                                                                                                                                                                                          SECREGATION. IT HAS RELAXATION OF SUPERCOILED DNA ACT PERFORMS THE DECATENATION EVENTS REQUIRED DURING THE OF A CIRCULAR DNA MOLECULE (BY SIMILARITY), SUBUNIT: COMPOSED OF TWO SUBUNITS: PARC AND PARE. SIMILARITY; BELONGS TO TYPE II TOPOISOMERASE FAMILY.
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                       AE001117; AAC66418.
L32861; AAC41408.1;
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                                                                       non-profit institutions as long as its content and this statement is not removed. Usage by an requires a license agreement (See http://www.isb-an email to license@isb-sjb.ch).
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87.5%;
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                                                                                      (See http://www.isb-sib.ch/announce/
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23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Borrelia.
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                                                                                                                                                                                                                                                                                                                     usual organization of disease spirochaetes
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Q04786;
01-NOV-1995
01-NOV-1995
30-MAY-2000
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SEQUENCE
                                                                                                                                                            DISULFID ACT_SITE
                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed
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                                                                                                                                                                                                                                      Hydrolase;
DISULFID
                                                                                                                                                                                                                                                                                                                                                              HSSP;
                                                                                                                                                                                                                                                                                                                                                                                  EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=93342067; PubMed=8341694; Somerville C.C., Colwell R.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A. STRAIN=ATCC 27562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BETA-N-ACETYLHEXOSAMINIDASE) (CHITOBIASE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00177; TOPOISOMERASE_II;
                                                                                                                                                                                                                DISULFID
                                                                                                                                                                                                                                                                                  Pfam; PF00728; Glyco_hydro_20; PRINTS; PR00738; GLHYDRLASE20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vulnificus: evidence for a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria; gamma subdivision; Vibrionaceae;
NCBI_TaxID=672;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vibrio vulnificus
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                                                                                                                                                                                                                                                                                                                                   InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nexosaminidases.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence analysis of the beta-N-acetylhexosaminidase rulnificus: evidence for a common evolutionary origin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CATALYTIC ACTIVITY: HYDROLYSIS OF TERMINAL NON-REDUCING N-ACETYL-D-HEXOSAMINE RESIDUES IN N-ACETYL-BETA-D-HEXOSAMINIDES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C. Natl. Acad. Sci. U.S.A. 90:6751-6755(1993).
FUNCTION: HYDROLYSIS OF TERMINAL, NON-REDUCING N-ACETYL-BETA-D-GLUCOSAMINE RESIDUES IN CHITOBIOSE AND HIGHER AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IN GLYCOPROTEINS.
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Q54468; 1QBB.
Similarity 6; Conserv
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                                                                                                                                                                 ; Glycosidase.
31 40
377 385
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519 519
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                                                                                                                   847 AA;
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(Rel. 32, Last seq
(Rel. 39, Last ann
  Conservative
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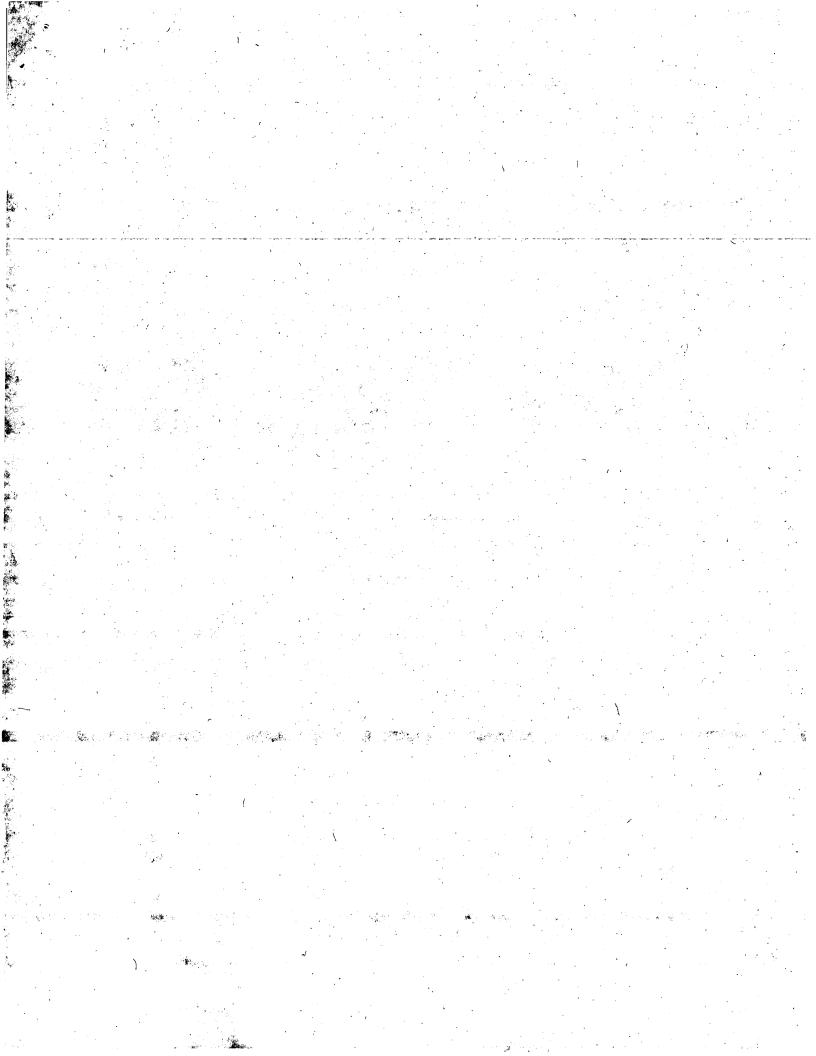
    Last sequence update)
    Last annotation update)
    (EC 3.2.1.52) (N-ACETYL-BETA-GLUCOSAMINIDASE)

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50.0%;
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ATP-binding.
mw; B5901F17B1CC7721 CRC64;
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CATALYTIC ACID (BINDS TO THE GILINKAGE) (BY SIMILARITY).
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                      Score 35; DB Pred. No. 43;
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Search completed: June 4, 2001, 12:24:05 Job time: 567 sec



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Minimum
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Perfect score:
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Listing first 45 summaries
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sp_bacteria:*
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sp_vertebrate:*
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13.062 Million cell updates/sec
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091nv4 arabidopsis
09vrh9 drosophila
09vrh8 drosophila
043546 lilium long
061818 mus musculu
09u1d0 leishmania
091uy1 arabidopsis
09kxc4 escherichia
                                                                                         Q13744 homo sapien
O77335 plasmodium
Q47307 escherichia
                                                                                                                           Q26654 sarcophaga
                                                                                                                                                                                 Q92663 homo sapie
Q92495 homo sapie
O58234 pyrococcus
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229 FYMGSKTLRGRN 240
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caenorhabdi	aquifex aeo	streptomyce			mycoplasma	clostridium	thermotoga	caenorhabdi	aspergillus	homo sapien	synechocyst	homo sapien		synechococc	s	9 molluscum c	clostridium	caenorhabdi	corynebacte	leishmania	0 arabidopsis	thermotoga	thermotoga	variovorax	bacteriopha

ALIGNMENTS

### 01-FEB-1997 01-FEB-1997 01-OCT-2000 MEDLINE-93055454; Put Porges A.J., Redecha Kimberly R.P.; Homo sapiens (Human). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 01-OCT-2000 (TrEMBLrel. FC GAMMA RECEPTOR I. SEQUENCE EMBL; L03418; AAA36049.1; HSSP; P12319; LALT. TISSUE-BLOOD; SEQUENCE FROM N.A. PFAM; PF00047; ig; PRODOM; PD002534; NCBI\_TaxID=9606; NTERPRO; IPR003006; -. . Clin Invest. Fc gamma receptor I family gene products in human mononuclear (TrEMBLrel. PRELIMINARY; 90:2102-2109(1992) PubMed=1430234; cha P.B., Doebele R., ; 1. 42632 MW; 02, 15, Created) Last sequence update) Last annotation update) PRT; D33D59398CEEA699 CRC64; 374 Pan L.C., B Salmon J.E.,

Conservative

0

Mismatches

Indels

0

Gaps

0

100.0%;

Score 64; Pred. No.

0.00026;

DB 4;

Length 374;

PRELIMINARY;

PRT;

375

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Best Local S
Matches 12
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01-AUG-1998
01-JUN-2000
Kawarabayasi Y., Sawada M., Horikawa H., Haikawa Y., Hino Y., Yamamoto S., Sekine M., Baba S.-I., Kosugi H., Hosoyama A., Naga Sakai M., Ogura K., Otsuka R., Nakazawa H., Takamiya M., Ohfuku Funahashi T., Tanaka T., Kudoh Y., Yamazaki J., Kushida N., Oguc Aoki K.-I., Yoshizawa T., Nakamura Y., Robb F.T., Horikoshi K.,
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058234;
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Ol-FEB-1997 (TrEMBLrel. 02, Last sequence update)
Ol-CCT-2000 (TrEMBLrel. 15, Last sequence update)
FC GAMMA RECEPTOR TYPE I (FC GAMMA RIB-FC GAMMA RECEPTOR)
                                                                                             Archaea; Euryarchaeota; NCBI_TaxID=53953;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL;
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01-FEB-1997
                                                   MEDLINE=98344137; PubMed=9679194;
                                                                STRAIN-OT3
                                                                                                                 Pyrococcus horikoshii
                                                                                                                             PH0498
                                                                                                                                     HYPOTHETICAL
                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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PFAM; PF00047; ig; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=93018827;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (DEC-1992) to the EMBL/GenBank/DDBJ databases
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                                                                           SEQUENCE
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                                                                        FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A.B.;
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(TrEMBLrel. 07, Last sequence up)
(TrEMBLrel. 14, Last annotation
L 44.0 KDA PROTEIN PH0498.
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                                                                                                                                                                                                                                                                                    .Conservative
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01-AUG-1998 (TrEMBLrel. 07, La
01-JUN-2000 (TrEMBLrel. 14, La
"HYPOTHETICAL 31.3 KDA PROPERTICAL 31.3 KDA
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01-AUG-1998
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      Ó9LPL9
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      09LPL9;
      01-OCT-2000 (TrEMBLrel. 15;

      01-OCT-2000 (TrEMBLrel. 15;
      15,

      01-OCT-2000 (TrEMBLrel. 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              thermophilic archaebacterium, DNA Res. 5:55-76(1998).
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae: Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II;
Brassicales; Brassicaceae; Arabidopsis.
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Hypothetical protein.
SEQUENCE 379 AA; 4
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                                                                                                                                   F2J10.16.
                                                                                                                                                                F2J10.16 PROTEIN.
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70.0%;
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1. 14, Last annotation updat
PROTEIN PH0394.
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Pred. No.
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Pred. No.
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horikoshii OT3.";
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Ohfuku Y.,
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                                                                                                      Q9Z2C5 PRELIMINARY;
Q9Z2C5;
01-MAY-1999 (TrEMBLrel. 1
01-MAY-1999 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
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Sakano H., Liu S.X., Yu G., Lee J., Lenz C., Pham P., Toriumi M.,
Chin C., Chlou J., Choi E., Chung M., Gonzalez A., Howng B., Liu A.,
Vaysberg M., Altafi H., Brooks S., Buehler E., Chao Q., Conn L.,
Conway A.B., Hansen N.F., Johnson-Hopson C., Khan S., Kim C., Lam B.
Miranda M., Nguyen M., Palm C., Shinn P., Southwick A., Davis R.W.,
Ecker J.R., Federspiel N.A., Theologis A.;
"The sequence of BAC F2J10 from Arabidopsis thaliana.chromosome 1.";
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AC015445; AAF76448.1;
"SEQUENCE 318 AA; 35231 MW; E722F2FBED9FD961 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                   Mus musculus (Mouse).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
                                                                                                                                                                                                                                                                                                                                                                   Yan Y., Li X., Wang A., Zhang G.;
"Molecular cloning and identification of full-length cDNA encoding high affinity Fc receptor for bovine IgG (Fc gamma RI).";
Vet. Immunol. Immunopathol. 75:151-159(2000).
BENDL; AF162866; AAF80477.1;
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                                                                                        MYOTUBULARIN
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SEQUENCE FROM N.A
                        NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                             Similarity 63.07; Conservative
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) (TrEMBLrel. 10,
) (TrEMBLrel. 13,
IN (EC 3.1.3.48).
                                                                                                                                                                                                                                                                                                                                              349 AA;
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                                                                                                                                                          PRELIMINARY;
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Rodentia;
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87.5%;
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15,
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Last annotation update)
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                                   Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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Kim C., Lam B.,
, Davis R.W.,
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SEQUENCE
                            Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
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"Characterization of the myotubularin from yeast to human.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-98409499; PubMed-9736772; Laporte J., Blondeau F., Buj-Bello A.,
                                                                                                                       ALL-1
                                                                                                                                                   ALL-1 PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PFAM; PF00372;
SEQUENCE 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=95181458; PubMed=7876233;
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Oestroidea;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sarcophaga peregrina (Flesh fly) (Boettcherisca peregrina).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INTERPRO;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Molecular cloning and sequencing of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=7386;
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Mol. Genet. 7:1703-1712(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      D29741; BAA06161.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IPR000387;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Metazoa; Arthropoda; Tracheata; Hexapoda; Neoptera; Endopterygota; Diptera; Brachyo; Sarcophagidae; Sarcophaga.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               72; hemocyanin;
1163 AA; 133401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                            Chordata;
Primates;
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                               Craniata; V
Catarrhini;
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fat body. Implications
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1.5e+02;
                                                              Vertebrata; Euteleostomi;
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Zweckbronner I., Beck J.D., Fey G.H.;
"MoLecular analysis of the chromosomal breakpoint and fusion
transcripts in the acute lymphoblastic SEM cell line with chromosomal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Exon/intron structure of the human ALL-1 translocations to chromosomal region 11q2
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Similarity
8; Conserv
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PS00354; HMGI_Y; UNKNOWN_1
PS50014; BROMODOMAIN_2; 1.
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Plasmid pUB2380.
Bacteria; Proteoba
Escherichia.
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01-MAY-2000
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                                                                                                                   origins for three different
submitted (AUG-1998) to the
EMBL; AJ008006; CAA07835.1;
Plasmid.
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01-NOV-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plasmodium falciparum
Eukaryota; Alveolata;
NCBI_TaxID=5833;
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01-MAY-1999
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                                                                                                   SEQUENCE
                                                                                                                                                      Albiger B., Bennett P.M., Commanducci A., Dodd H.M., Lett M.C., "pUB2380: a ColD-like resitance plasmid with three replication origins for three different functions."; Submitted (AUG-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
TRANSPOSON=IS1294;
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FIFRO01064; -
FIFRO0125; CRYSTALLIN_BETAGAMMA; UNKNOWN.
FS00225; CRYSTAL
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    score 36; DB; Pred. No. 64; 0; Mismatches
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Last annotation update)
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20

2 YMGSKTLRGRN

237

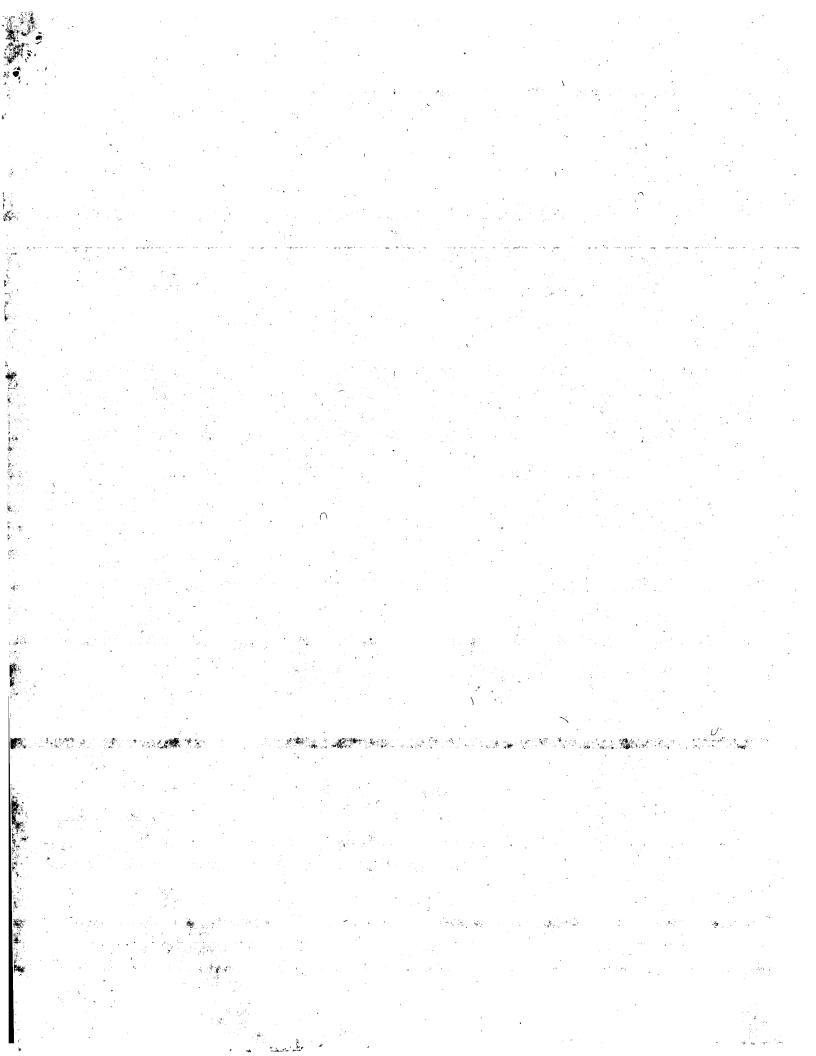
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Q9LNV4;
01-OCT-2000
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                                                                                                         Q9VRH9
                                                                                                                        Q9VRH9
                                                                                                                                                                                                                                                                                                                                                                     Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases EMBL; AC022464; AAF79558.1; - SEQUENCE 415 AA; 48371 MW; 8C86C784BOAC2C68 CRC64;
Drosophila melanogaster (Fruit fly)
                                    SLGA PROTEIN
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Magnoliophyta; eudicotyledons; core euc
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Lee J., Lenz C., Li J., Liu A., Liu J.,
Yguyen M., Palm C., Pham P., Sakano H.,
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                                                                                                                                                                                                                                                2 YMGSKTLRGRN 12
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01-MAY-2000
01-JUN-2000
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George R.A., Lewis S.E., Richards S., Ashburner M.,
Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., (
                            Drosophila
                                                                                    SLGA PROTEIN
                                                                                                                                                                                                                         Q9VRH8
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Liu X., M
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Jalali M.,
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     Eukaryota;
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491 FYFGAKLVRG 500
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AE003568; AAF50819:1;
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Similarity 60.0%;
6; Conservative
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  melanogaster (Fruit
Metazoa; Arthropoda;
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Drosophilidae; Drosophila.
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AA; 75951 MW;
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era; Muscomorpha;
        Insecta;
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LIMY.
LIMY Iongiflorum (Trumpet lily).
Lilium longiflorum (Trumpet lily).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Eukaryota; Liliopsida; Liliales; Liliaceae; Lilium.
Magnoliophyta; Liliopsida; Liliales; Liliaceae; Lilium.
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                                                                                                                                                                                                                          SERINE PROTEINASE (FRAGMENT).
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"The genome sequence of Drosophila melanogaster.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Science 287:2185-2195(2000).
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Sutton G.G., Wortma
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INTERPRO; IPR002872; -
PFAM; PF01619; Pro_dh; 1.
SEQUENCE 671 AA; 76113 MW;
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Ephydroidea; Drosophilidae; Drosophila.
NCBI_TaxID=7227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             493
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S.E., Richards S., Ashburner M., Henderson S.N.,
an J.R., Yandell M.D., Zhang Q., Chen L.X.,
rs Y.-H.C., Blazej R.G., Champe M., Pfelifer B.D.,
Baxter E.G., Helt G., Nelson C.R., Miklos C. I. A., An H.-J., Andrews-Pfenci
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60.0%;
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Pred. No. 1.3e+02;
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SEQUENCE
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MENDEL; 11876; Lillo;1086;11876.
INTERPRO; IPRO00209; -.
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                                                                                                                                                                                                                                                                                              microsporocytes.
                                                                                                                                                                                                                                                                                                                                         STRAIN=CV. HINOMOTO; TISSUE-FLORAL BUD; MEDLINE=96051386; PubMed=7584025;
                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=CV. HINOMOTO; TISSUE=FLORAL BUD;
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[1]
                                                                                                                                                                            PFAM; PF00082; Peptidase_S8; 2.
PRINTS; PR00723; SUBTILISIN.
PROSITE; PS00136; SUBTILASE_ASP;
PROSITE; PS00138; SUBTILASE_SER;
NON TETT
                                                                                                                                                                                                                                                                                    DNA Res. 1:15-26(1994).
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                                                        YYSGHTLRGRN 515
                                                                             YMGSKTLRGRN 12
                                                                                                  Similarity 63.07; Conservative
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                                                                                                                                                           813 AA;
           June
                                                                                                                                                                                                                                                                                                        of cDNAs induced in meiotic prophase in lily
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          2001, 12:23:26
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                                                                                                            Score 36;
Pred. No.
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1.6e+02;
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum DB
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/SIDS6/gcgdata/geneseq/geneseqp/AA1997.DAT:
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                                                                                                                                                                                                                                                                                                               /SIDS6/gcgdata/geneseq/geneseqp/AA2000.
/SIDS6/gcgdata/geneseq/geneseqp/AA2001.
                                                                                                                                                                                                                                                                                                                                                         /SIDS6/gcgdata/geneseq/geneseqp/AA1999
                                                                                                                                                                                                                                                                                                                                                                          'SIDS6/gcgdata/geneseq/geneseqp/AA19
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                                                                                                                          DВ
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                                                           W60557
Y33183
¥96183
                   R91439
                                         R22549
                                                                                                                                                                                     SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /geneseqp/AA1988.DAT:*
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	31	31	31	32	32	32	32	3 2	32	32	32	32	32	32	ω ω	33	<b>ω</b>	33	ω 3	<b>ω</b>	ω	ω G	34	36	36	φ 9	36	36	36	57	57	57	57	57
	54.4	54.4	54.4	56:1	56.1	56.1	56.1	56.1	56.1	56.1	56.1	56.1	56.1	56.1	57.9	57.9	57.9	7.	57.9	57.9	57.9	7.	9.	63.2	ω.	ú	ω ·	ω ·	٠	•		100.0		100.0
	255	185	58	992	969	802	623	525	506	451	428	284	283	25	1165	749	696	696	395	371	353	193	88	2004	2004	2004	2004	1879	410	399	377	374	374	374
	21	21	21	21	21	21	15	20	21	21	18	21							20	21	21	18	20	21	20	19	17	14	12	21	20	21	21	19
	Y74710	Y58587	G00321	G33920	G33921	G33922	R49578	Y25881	Y94986	G39206	W11883	G39207	G39208	Y25887	R37309	P70286	Y17207	W55698	Y31732	B54247	Y67867	W55378	Y35386	в10469	W89402	W37052	R99464	R42455	R12428	B43683	W86195	Y96226	Y96134	W97834
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## ALIGNMENTS

RESULT W60557 W60557; W60557 standard; peptide; 12 AA. Synthetic. immunohistochemical;
cell transformation; Oligopeptide from extracellular domain of CD64 De Kruif CA, (UYUT-) RIJKSUNIV UTRECHT. 08-OCT-1996; 07-OCT-1997; W09815833-A1. Extracellular domain; 18-AUG-1998 16-APR-1998 (first entry) Logtenberg 96EP-0202791 97WO-NL00557 mutation; immunofluorescent analysis; detection; CD64; identification; antibody; oncogene.

57 57 57 57 57 57 57 57 57 57 57 57 57 100 00 W00859 R91438 Oligopeptide from Human sFC-gammaRI Human macrophage-s Human FCRI (CDNA c Human macrophage-s Human macrophage-s Human Human Human Human Human n macrophage-s n FcRI (cDNA c n FcRI (cDNA c receptor receptor

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies WPI; 1998-240964/21.

Synthetic oligopeptides W60537-61 are derived from the extracellular

25 polyethylene rods as

isolated

Example 1; Page 29; 40pp; English

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Best Local
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                                                                                                                      Baell JB,
Maxwell KF,
                                                                                                                                                                                                             11-SEP-1998;
06-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                      atomic coordinate; bioactive compound design; computer-assisted; drug design; therapy; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; IgE; tissue damage; hypersensitivity; inflammatory cell recruitment;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     domain of CD64. They were synthesised on 25 polyethylene rods adjacent, non-overlapping sets. Peptide specific phages were affinity binding to the CD64-covered rods. The specification
                                                                                                     Powell MS;
                                                                                                                                                                        (ILEX-) ILEXUS PTY LTD.
                                                                                                                                                                                                                                                                 04-FEB-1999;
                                                                                                                                                                                                                                                                                                                                      W09940117-A1
                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                          anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fc receptor; Fc-gammaRIIa; human; FCR; model; three-dimension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Y33183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Y33183 standard; Protein; 261 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method for the identification of a peptide able to bind specifically to
                                                                                                                                                                                                                                                                                                   12-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human sFc-gammaRI protein fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                             inflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100 nes 12; Conservative
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                                                                                                                  Epa V, Garrett, McCarthy TD,
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                                                                                                                                                                                                                                                                                                                                                                                                                             modulator; Fc-gammaRIa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                           98US-0099994.
98US-0073972.
                                                                                                                                                                                                                                                                 99WO-IB00367.
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                                                                                                                                                                                                                                                                                                                                                                                                          immunoprotective;
                                                                                                                  TPJ, Hogarth PM,
McKenzie IFC, Pi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 57;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                       immune function sFc-gammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 19;
                                                                                                                 Pietersz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The specification describes
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                                                                                                                                     Matthews
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                                                                                                                                                                                                                                                                                                                                                                                                                       regulation;
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Three-dimensional structures computer assisted drug design

structures and models of Fc receptors, useful in

This amino acid sequence was predicted from the cDNA sequence of CDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25, p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90

Example 10; Page 94a; 160pp; English

WPI; 1999-539978/45

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RESULT
R22549
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Best Local
                                                                                                                                                                   N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIA and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to chemical compounds can be designed to block binding of immunoglobulin to
                                                                                                             New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                   WPI; 1992-056864/07
N-PSDB; Q21179.
                                                                                                                                                                                                                    Seed B,
                                                                                                                                                                                                                                                                                        13-JUL-1990;
                                                                                                                                                                                                                                                                                                                       15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                       23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                         W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R22549 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               an RC receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of
                                                                                                                                                                                                                                                      (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human macrophage-specific FcRI receptor encoded by clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  oligonucleotides, peptides, peptidomimetics and small organic molecumby computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to reduce IGG- or IGE-mediated tissue damage, e.g. IGG- or IGE-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the FcR can be used to design bioactive chemical compounds, oligonucleotides, peptides, peptidomimetics and small organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; Page 316-317; 326pp; English
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12; Conser
                                                                                                                                                                                                                      Aruffo
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                                                                                                                                                                                                                      Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              344
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Pred. No. 0.00078;
Pred. No. 0.00078;
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RESULT
R91439
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Matches 12
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                      The amino acid sequence (R91439) of human FCRI was detd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other CDNA clones, pl35 (N14717) and p90 (N14719), coded for a variants of the FCRI sequence; the C-terminal sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to
                                                                                                                                                                                                                                                                   Cloning of cDNA encoding cell surface of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                    Aruffo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human FcRI (cDNA clone p98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-OCT-1996
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                                                                                                                                                                                                                Example 10; Column 55-56; 79pp; English.
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les 12; Conserv
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DB; T14718.
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88US-0160416.
89US-0379076.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            s-0553759.
s-0139273.
is truncated compared
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "amino acid 58 is Leu in 
'translated product"
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Best Local 9
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Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                 panning; immunodiagnosis; diagnosis; immunotherapy; gene th
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human macrophage-specific FcRI.
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1 Similarity 100.
12; Conservative
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117
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                                                                                                                                                                                                                                                                TCC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  . A fusion protein in increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy;
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Query Match
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25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
                                                                                                                                                                                                                                                                                                         surfaces other than macrophages,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the invention designed to isolate CSA nucleic acids. The method based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of
                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                          protein of FcRI and
                                                                                                                                                                                                                                                                                                                                                   ability to
                                                                                                                                                                                                                                                                                                                                                                                                               sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              animals, including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proteins they encode, are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           physical selection of cells expressing the antigen by adhesion to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-586382/55.
N-PSDB; A50631.
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                                                                                                                                                                                                                                                                                                                              immune complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of 19G, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on the contract of the cell surfaces of the contract of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cell surfaces of the cel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of immune-mediated infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Column 53-55; 75pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mmunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                          potencies
ch 100.0%;
l Similarity 100.0%;
12; Conservative 0
                                                                                                                                                                           344 AA;
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90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                89US-0379076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92US-0983647
88US-0160416
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338
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306
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305
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268
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220
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216
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                                                                                                                                                                                                                                                                          a receptor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 These disorders
       0;
                                                                                                                                                                                                                                          in
   Score 57; DB 21;
Pred. No. 0.0011;
; Mismatches 0;
                                                                                                                                                                                                                                es, is useful c ligand will be therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGT"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diseases, and disorders in
                                                                                                                                                                                                                                                                      in therapy. A fusion be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              include asthma,
                                                               Length 344;
Indels ·
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The method is
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    parasitic
Gaps
0;
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1 TSSEYQILTARR 12

21-MAY-1992

(first entry)

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X A X A X E
                                                                                                                                                                                                            RESULT
R22550
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R20811
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XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six optential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, p135 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the
                                                                                                       R22550;
                                                                                                                                                                       R22550 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N-PSDB; Q21178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rapid immunoselection cloning technique; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R20811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R20811 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                     241 tsseyqiltarr
                                                                                                                                                                                                                                                                                                                                                                                          1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ch 100.0%;
1 Similarity 100.0%;
12; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90WO-US04986
                                                                                                                                                                                                                                                                                                                                                     252
                                                                                                                                                                          374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 57; DB 13; Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cell surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neoplasms, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               it -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     s. A fusion protein
increase the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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Human macrophage-specific FcRI receptor encoded by clone p90.

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                          FIFXXXXXXX
                                                                                                                                                                                                                                                                                                                                                                 Query Match 100
Best Local Similarity 100
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p98/x2 (see Q21179) both predict Leu. sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seed B, . Aruffo A, Amiot M,
                                                                                                         Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FCRI; Fc receptor.
                                                                                                                                                    Human FcRI (cDNA clone p135 product).
                                                                                                                                                                               30-OCT-1996 (first entry)
                                                                                                                                                                                                           R91438;
                                                                                                                                                                                                                                       R91438 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This amino acid sequence was predicted from the cDNA sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
Misc-difference
                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                         Misc-difference
                                                                                                                                                                                                                                                                                                            241
                                                                                                                                                                                                                                                                                                                         1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                   8
                                                                                                                                                                                                                                                                                                         tsseyqiltarr 252
                                                                                                                                                                                                                                                                                                                                                                                                                                      374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90US-0553759
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                                         Location/Qualifiers 25
 58
                          /note-
                                                                                                                                                                                                                                                                                                                                                                              100.0%;
           "amino acid 25 is Thr in clone p90 and p98 translated products"
                                                                                                                                                                                                                                       374 AA
                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                 Score 57; DB 13;
Pred. No. 0.0012;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                          Length 374;
                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                 0,
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Best Local Similarity 100.0%;
Matches 12; Conservative 0
  01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (MO0859) of the FCRI sequence, and a third clone, p9 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5506126-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1996
                                                                                                                       09-APR-1996
                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                            therapy; diagnosis;
                                                                                                                                                                                                                                                                    Cell surface antigen; cloning; immunoselection; immunotherapy;
                                                                                                                                                                                                                                                                                                          Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                                                                     30-0CT-1996
                                                                                                                                                                                                                                                                                                                                                                                             W00859;
                                                                                                                                                                                                                                                                                                                                                                                                                                W00859 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The amino acid sequence (R91438) of human FCRI was detd. cDNA clone, p135 (T14717), obtd. from a cDNA library usi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; T14717.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-FEB-1988;
                                                                                   25-FEB-1988;
                                                                                                                                                                  US5506126-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         374 AA;
                                                                                                                                                                                                                                                                                                                                                     (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92US-0983647.
88US-0160416.
89US-0379076.
89US-0553759.
93US-0139273.
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                                                                                   88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "amino acid 58 is Val in p90 clone
    translated product"
                                                                                                                                                                                                                                                 vector; FcRI; Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                    374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 57;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The amino acid sequence (W00859) of human FCRI was detd. from a cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of the FC portion of IgG, normally located on cell surfaces of the FC portion of IgG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FcRI sequence. A fusion protein of FcRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-200279/20.
N-PSDB; T14719.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                          cloning
                                                                                                                                                                                                                                                                                                                                                                                      07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                  W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                          W80448 standard; Protein; 374 AA.
                                                                                                                             Misc-difference
                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                   Fc receptor I; FcRI;
                                                                                                                                                                                                                                                                                                                                                              Human Fc receptor
Misc-difference
                         Misc-difference
                                                                             Misc-difference
                                                                                                      Misc-difference
                                                                                                                                                                                 Misc-difference
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nilarity 100.0%;
Conservative 0
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93US-0139273.
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45
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141
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103
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                                                                                                                                                                                                                       'note=
                                                                                                                                                                                                                                                                                                                                       cell surface antigen; lymphocyte; human;
                                                                                                                                           "encoded by
                                                                                                                                                                   "encoded by CTG"
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Pred. No. 0.0012;
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                                     by ccc"
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                                                                                                                                            ccc,"
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                                                                                          TCC
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                GGC"
                                                                  CAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 374;
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W97833 ID W97 XX AC W97

W97833 standard;

Protein; 374 AA:

1

W97833;

밁 Qy

241 tsseyqiltarr

252

1 TSSEYQILTARR 12

Query Match Best Local S Matches 12

l similarity 100 12; Conservative

0;

100.0%;

Score 57; DB 19; Pred. No. 0.0012; ; Mismatches 0;

Length 374; Indels

0;

Gaps

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XX
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13-JUL-1990;
21-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                           This is deduced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                            adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                             expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                            were also obtained.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            human lymphocyte antigenic sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB;
Sequence
                                                                                                                                                                                                                                                                                                                                        from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see x07372) and p98 (see x07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                          treatment of immune-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                    immunoselection cloning method from a cDNA library expressed
                                                                             .mmunotherapeutic applications,
                                                                                                                                                                                                                                                                                                                                                                                            The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                           the amino acid sequence of human Fc receptor I (FCRI), as from cDNA clone p135 (see V63456) isolated using a rapid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10;
                                       including
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374 AA;
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89US-0379076.
90US-0553759.
97US-0861205.
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176
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256
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                                                                 infections,
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This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p90 (see X07372) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see X63456) and p98 (see X07373)
                                                                                                                                                                                                                                                                                                   01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                                                                                       New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
             induction therapy.
were also obtained
                                                                                                                Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                             WPI; 1998-609251/51.
N-PSDB; X07372.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human Fc receptor I.
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89US-0379076
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103
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99
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176
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expression of an antigen
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            from mammalian
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Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in eukaryotic cells and selection of cells expressing the antigen adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                 /note=
Misc-difference 141
                                                                                                                                                                                                                                     Misc-difference 45
                                                                                                                                                                                                                                                                                                                                                                                  Human Fc receptor I
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                                                                                                                                                                                                                                                                                                                                                                                                                            W97834;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                         Misc-difference
                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                           Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunotherapeutic applications, including
treatment of immune-mediated infections,
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159
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176
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/note= ? 64

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CAC" ACC" CGG. ACC" Location/Qualifiers

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                                                                                                                                                                                                                                                                                                                                                C from a single patient undergoing extracorporeal interleukin-2
C induction therapy. Clones pl35 (see V63456) and p90 (see X07372)
C were also obtained. A novel method for cloning cDNAs from mammalian
C expression libraries is based on transient expression of an antigen
C in eukaryotic cells and selection of cells expression the antigen by
C adhesion to an antibody coated substrate. The method is useful for
C the isolation and cloning of any protein which can be expressed and
C transported to the cell surface membrane of a eukaryotic cell. It
C has been used to clone genes (see V63442-63) encoding cell surface
C antigens from mammalian lymphocytes (see W80440-55). The isolated
C genes can be expressed in a prokaryotic or eukaryotic host cells to
C produce the encoded protein. The invention also provides high
C efficiency expression vectors (see V63441 and V63444) which allow
C the generation of very large mammalian expression libraries. The
C purified genes and proteins are useful for immunodiagnostic and
C purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
            panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy;
immune disorder; infection; asthma; immune-complex disease;
                                       Macrophage; FcRI; cell surface antigen; human; immunoselection;
                                                                    Human macrophage-specific FcRI.
                                                                                                19-DEC-2000
                                                                                                                                                     Y96134 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone 998/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in Coells. The CDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1998-609251/51.
N-PSDB; X07373.
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                                                                                                                                                                                                                                                                                                                                                                                                mmunotherapeutic applications,
                                                                                                                                                                                                                                    1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                        including
                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                       immune-mediated
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88US-0160416.
89US-0379076.
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97US-0861205
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                                                                                                                                                     Protein;
                                                                                                                                                                                                                                                                                                                                                                          humans
                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORP.
                                                                                                                                                     374
                                                                                                                                                                                                                                                                            0;
                                                                                                                                                     AA.
                                                                                                                                                                                                                                                                                      Score 57; DB 19;
Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                               including the diagnosis and
                                                                                                                                                                                                                                                                            Mismatches
parasitic
                                                                                                                                                                                                                                                                                                                                                                                     diseases,
disease;
                                                                                                                                                                                                                                                                                                     19;
                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                     Length 374;
                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                     and disorders of
Fc receptor
                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                          Gaps
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25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
           Stamenkovic I,
                                                                                                                      01-DEC-1992;
                                                                                                                                                 28-OCT-1998;
                                                                                                                                                                           29-AUG-2000.
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                                     (GEHO ) GEN HOSPITAL CORP
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/note= ; 216

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CAA" AAT"

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88US-0160416. 89US-0379076. 90US-0498809. 90US-0553759.

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9205-0983647 98US-0181612

WPI; 2000-586382/55 N-PSDB; A50592.

physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,

diseases, and disorders in

immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FcRI and a receptor ligand will be helpful to increase

The present sequence is that of a human macrophage specific as deduced from cDNA clone pl35 (see A50592), which was isol

as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of

based upon transient expression of a CSA in eukaryotic cells and

the invention designed to isolate CSA nucleic acids.

The method is

of.

Example 10; Column 53-55; 75pp; English.

Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic

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В
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                                                                                                                                                                                                                                                                         ¬EP1006183-A1
                                                                                                                                                                                                                                                                                                                                                                                            Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IgG; immunosuppressive; antirheumatic; cytostatic; antiinflammatorv: AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-SEP-2000 (first entry)
                                                                                                                        03-DEC-1998;
                                                                                                                                                                                                                          07-JUN-2000
                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                        systemic lupus erythematosus; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Y96226 standard; Protein; 374 AA
N-PSDB; A27466
                         WPI; 2000-367968/32
                                                                                                                                                                         03-DEC-1998;
                                                                     (PLAC ) MAX PLANCK GES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tsseyqiltarr 252
                                                                                                                        98EP-0122969
                                                                                                                                                                       98EP-0122969
                                                                        FOERDERUNG WISSENSCHAFTEN.
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RESULT 15
W86195
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local :
                                                           13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      may be modified to produce recombinant versions. The recombinant FC receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays t determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W86195 standard; Protein; 377 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is the imman interpretary an important regammaRI is also known as CD64, receptors play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds opsonised by serum immunoglobulins. The resulting complex then binds opposite the process of the process of the process of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of the present of 
                                                                                                                                                                                  01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                             07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                        15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                 US5849898-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human Fc receptor I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W86195;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pathways are activated, leading to immune response. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is the human high affinity Fc receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rheumatoid arthritis
(GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 TSSEYQILTARR 12
||||||||||
| 241 tsseyqiltarr 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                   92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CD36; cell surface antigen; human; cDNA library
                                                              95US-0485447
                                                                                                                          90US-0498809
                                                                                                                                                                                                                                                                             95US-0485447
                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note= "this sequence contains a considerable number of differences from the sequence deduced from the DNA sequence given in the specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 57;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and
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0

Matches 12; Query Match Best Local

Conservative

0;

1 TSSEYQILTARR 12

Local Similarity

100.0%;

Score 57; DB 21; Pred. No. 0.0012; Mismatches

DB 21;

Length 374; Indels.

0;

Gaps

0

Sequence

374 AA;

the potencies of antibodies in therapy.

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Search completed: June Job time: 203 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This polypeptide human Fc receptor I (FCRI). FCRI cDNAs (see V81213) were isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extreorporeal interleukin-2 induction therapy. DNA sequence analysis revealed that the cDNAs encoded type I integral membrane proteins with 3 extracellular immunoglobulin domains. The invention provides a novel method for cloning cDNAs from mammalian expression libraries. This is based on transient expression of an antigen in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and molecular cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell.
                                                                                                                                                                                                                                                                  Query Match 100.0%; 9
Best Local Similarity 100.0%; 1
Matches 12; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA encoding human CD40 antigen - useful for cloning cDNA encoding cell surface antigens, constructing cDNA libraries, expression vectors for expression in eukaryotic cells or their fragments
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CD40_cDNA (see V81198) is specifically claimed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; V81213.
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Oquendo C,
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244 tsseyqiltarr 255
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Seed B,
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Pred. No. 0.0012;
); Mismatches 0;
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), Stamenkovic I, Stengelin S;
                                                                                                                                                                                                                                                                                                                           Length 377;
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Result
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Maximum DB
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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57
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Gapop 10.0 , Gapext 0:5
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
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    PCT US95-16311-1
US-08-469-486-52
US-08-469-658-52
US-09-036-987a-7
US-08-986-217-3
US-09-154-874-9
US-09-170-670-5
US-09-183-412-5
US-09-183-412-5
US-08-683-838A-13
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US-08-375-799-15
US-08-375-799-15
US-09-090-793-9
US-09-032-372-1
US-08-240-3572-1
US-09-335-409-16
US-08-372-652-2
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US-09-469-721-2
PCT-US95-17023-2
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US-08-372-652-1
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US-09-198-484-2	5264554-2	US-08-478-208-32	PCT-US95-10426-34	PCT-US94-01553A-34	US-08-985-659-37	US-08-914-679A-5	US-08-890-383-5	US-08-704-706A-36	US-08-468-698-34	US-08-468-220-34	US-08-645-971-4	US-08-468-700-36	US-08-683-838A-4	US-08-600-908A-4	US-08-687-399-4	US-08-343-804-4	US-08-459-610-4	
Sequence 2, Appli	Patent No. 5264554	Sequence 32, Appl	Sequence 34, Appl	Sequence 34, Appl	Sequence 37, Appl	Sequence 5, Appli	Sequence 5, Appli	•	Sequence 34, Appl	Sequence 34, Appl	Sequence 4, Appli	Sequence 36, Appl	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 4, Appli	

### ALIGNMENTS

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US-08-332-562A-94
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                                                                                           INFORMATION FOR SEQ ID NO:
               SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                 REFERENCE/DOCKET NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 54;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
TELEFAX: 004126
                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                   NAME: BENT, Stephen A. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                          APPLICATION NUMBER: US 07/896,457 FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3000 K Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                       Patentin Release #1.0, Version #1.30
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ENTION: FC RECEPTOR FOR IMMUNOGLOBULIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HOGARTH,
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linear
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                                                                                                                                                                                                                                                                                                                                31-OCT-1994
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                                                                                            94:
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Query Match

63.2%; Score/36;

DB 2;

Length 91;

66.7%;

No.

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RESULT 3
US-08-752-929-15
; Sequence 15, Ap
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US-08-375-709-15
Sequence 15, Application US/08752929 Patent No. 5798259
                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                     Query Match
                                                                                                                                                         Matches
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Matches 8; Conserv
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Sequence 15, Approx
80, 568389
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                                                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: JP 4-147945
FILING DATE: 15-MAY-1992 -
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 14-MAY-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Release #1.0, Version #1.30 SURRENT APPLICATION DATA:
                                                                                          905 TASEQQALTAR 915
                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ITLE OF INVENTION: Synthesizing Enzym
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPLICANT: KATO, Seishi
PPLICANT: KONDO, Kiyosi
ITLE OF INVENTION: Gene
                                                                                                                                                                                                                                                                                                                                                               TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: WEGNER, Harold C.
REGISTRATION NUMBER: 25,258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: FILING DATE: 14-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                        1 TSSEYQILTAR 11
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                                                                                                                                                       Conservative
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                                                                                                                                                                        63.2%;
72.7%;
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Synthesizing Enzymes and Process for Production of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US 08/178,251
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                                                                                                                                                                        Score 36; DB Pred. No. 64;
                                                                                                                                                         Mismatches
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                                                                                                                                                                                       Length 2004;
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US-09-090-793-9
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                                                                                                                                                                     Sequence 9, Application No. 6140486
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                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
CURRENT APPLICATION NUMBER: US/09/09,793
CURRENT FILING DATE: 1998-06-04
EARLIER APPLICATION NUMBER: 60/048,650
EARLIER FILING DATE: 1997-06-04
                                                                  APPLICANT: Calgene, LLC
TITLE OF INVENTION: Production of polyunsaturated fatty acids by expression
TITLE OF INVENTION: of polyketide-like synthesis genes in plants
FILE REFERENCE: CGNE.131.0105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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APPLICATION NUMBER: US 08/375,709
FILING DATE: 20-JAN-1995
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: Foley's Lardner
STREET: 3000 K Street, N.W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 53
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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905 TASEQQALTAR 915
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 0 FILING DATE: 14-MAY-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER:
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FILING DATE: 15-MAY-199;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2004 amino acids
                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KATO, Seishi
KONDO, Kiyosi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YAMADA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-NOV-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kazunaga
                                                                                                                                                                                                                                                                                                                                                                                          63.2%; Score 36; DB 72.7%; Pred. No. 64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymes and Acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP 4-147945
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                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 2004;
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Query Match
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; ORGANISM: Shewanella putrefaciens
US-09-090-793-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn Ver. 3
SEQ ID NO 9
                                                                                 STRANL.
TOPOLOGY: LA..
IMMEDIATE SOURCE:
LIBRARY: SYNOR/
ONE: 78191
                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                         NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH: 2004
                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94304
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TLE OF INVENTION: CELL CYCLE RELATED PROTEINS MBER OF SEQUENCES: 13
                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 TSSEYQILTAR 11
                                                                                                                                                                               ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Palo Alto
                                                                                                                                                               amino acid
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                                                                                                                                                                            395 amino acids
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                                                                                                          SYNORAB01
                                                                                                                                                                                                                                      650-845-4166
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lal, Preeti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hillman,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Corley, Neil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yue, Henry
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                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM Compatible
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                                                                                                                                                   single
                          57.9%;
70.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    for Windows Version 2.0
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          Score 33; DB 3; Pred. No. 40; 2; Mismatches
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Pred. No.
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                                    Length 395;
          Indels
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       Gaps
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Query Match
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                                                                                                                                                                                                                Sequence 16, Application US/09335409 Patent No. 6121029
                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Cyr, Devon
APPLICANT: Goetlach, Joein
TITLE OF INVENTION: GENES FOR THE BIOSYNTHESIS OF EPOTHILONES
FILE REFERENCE: 4-30582A
CURRENT APPLICATION NUMBER: US/09/335,409
CURRENT FILING DATE: 1999-06-17
NUMBER OF SEQ ID NOS: 30
                                                                                                                             APPLICANT: Molnar, Istvan
APPLICANT: Zirkle, Ross
                                                                                                                                                              APPLICANT: Schupp, Thom APPLICANT: Ligon, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tent No.
                                                                                                                                                                                                                                                                                                                   1126 TTDLYQVLAAKR 1137
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/240,357
FILING DATE: 10-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Gordon, Alan M. REGISTRATION NUMBER: 30 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                  1 TSSEYQILTARR 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ELEPHONE:
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5. 5578481
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
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                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         201-831-3244
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , Alan M.
, Alan M.
30,637
                                                                                                                                                                                  Thomas
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Cardiac Adenylyl Cyclase
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                    Length 1165;
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; TYPE: PRT ; ORGANISM: Sorangium cellulosum US-09-335-409-16

SEQ ID NO 16. LENGTH: 185

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Query Match
Best Local Similarity
Thes 7; Conserv
                             PCT-US95-16311-2
                                             RESULT
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Sequence 2, Application PC/TUS9516311 GENERAL INFORMATION:
                                                                                                                                                                       Query Match
Best Local S
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Patent No. 5932699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                          NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 451 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/372,652
FILING DATE: 13-JAN-1995
                                                                                          357 TQEEYALLTA 366
                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 13-JAN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 617/542-8906
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APPLICANT: Choi, Hueng-Sik
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                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 02110-2804
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STATE: MA
                                                                                                                                                                                                                                                                   TOPOLOGY:
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TELEX: 20015
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                                                                                                                         TSSEYQILTA 10
                                                                                                                                                      l Similarity
6; Conserv
                                                                                                                                                                                                                                                                                                amino acid
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225 Franklin Street, Suit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Moore, David
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                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                              not relevant
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Pred. No. 1.2e+02
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Pred. No. 43;
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                                                                                                                                                        Mismatches
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                                                                                                                                                                                    Length 451;
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; MOLECULE TYPE::
PCT-US95-16311-2
                                                                                                                                                                                                                                                                                     Sequence 2, Application Patent No. 6005086 GENERAL INFORMATION:
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Matches 6; Conserv
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APPLIANT NUMBER: 13-JAN-1995
FILING DATE: 13-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T:
NAME: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                               TITLE OF INVENTION: ME
TITLE OF INVENTION: BY
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                        1 TSSEYQILTA 10
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                                                                                                                                                                                                                                                                                                                                                                                                        357 TQEEYALLTA 366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                   STREET: 444 South
CITY: Los Angeles
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Fish & KICHELLOCK STREET: 225 Franklin Street, Suite 3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                COUNTRY:
                                                                                                                               ADDRESSEE: Pretty, Schroeder, Brueggemann & Clark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 451 amino acids
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                                                                    90071
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                                                                                                                                                                                                                                                                                                                        Application US/08372183
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                                                                                                                                                                                                  Evans, Ronald M.
Forman, Barry M.
Weinberger, Cary A.
VENTION: METHOD FOR MODULATING PROCESSES MEDIATED
VENTION: BY FARNESOID ACTIVATED RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         not relevant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       54.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08/372,652
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Pred. No. 1.2e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 451;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:

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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      equence 2, Application US/09469721 atent No. 6184353
                                                            TELEFAX: 619-546-9392
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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SEQUENCE CHARACTERISTICS:
                                                                                                                                REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein -372-183-2
                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Weinberger, FITLE OF INVENTION: METITLE OF INVENTION: BY
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 OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Reiter, Stephen E. REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                 NAME: Reiter, Stephen REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 469 amino acids TYPE: amino acid TOPOLOGY: linear
                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                    SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
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                                              ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                        9007
                               amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Los Angeles
                                                                                                                                                                                                                                                                                                                                                                                                                                                       CA
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                                              469 amino acids
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619-546-9392
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                                                                                                                619-546-4737
protein
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METHOD FOR MODULATING PROCESSES MEDIATED
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                                                                                                                                                                                                                                                                                                   US/09/469,721
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Pred. No.
                                                                                                                                                 9844
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9844
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Best Local Similarity
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                                                                                                                              US-08-496-631-2
                                                                                                                                             RESULT
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Best Local Similarity
Matches 6; Conserv
                                                                                              Sequence 2, Applic Patent No. 5728548
                                                                              GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                            375 TQEEYALLTA 384
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 469 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                              APPLICANT: Bowman, Michael TITLE OF INVENTION: STEROI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Relier, Stephen E.
REGISTRATION NUMBER: 91,192
REFERENCE/DOCKET NUMBER: 94;
                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORRESPONDENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U
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CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER:
                                                                                                                                                                                                                          1 TSSEYQILTA 10
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                                                                                                            Application US/08496631
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VENTION: METHOD FOR MODULATING PROCESSES MEDIATED

VENTION: BY FARNESOID ACTIVATED RECEPTORS
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                                                                                                                                                                                                                                                                                                                                                                       linear
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Genetics Institute,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
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                                              STEROID RECEPTOR RR1
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Pred. No.
                                                                                                                                                                                                                                                                          Score 31;
Pred. No. 1
                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                       Length 469;
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                                                                                                                                                                                                                                                           Indels
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STREET: o, Cambridge

87 CambridgePark Drive

Massachusetts

COUNTRY:

U.S.A.

02140

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Query Match
Best Local Similarity
Ches 6; Conserva
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US-08-372-652-1
                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/372,652
FILING DATE: 13-TAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00786/246001
TELECOMMUNICATION INFORMATION:
THE LECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein -08-496-631-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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REGISTRATION NUMBER: 32,724
REFERENCE,DOCKET NUMBER: GI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
TELETANA: 4617, 498-8224
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
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COUNTRY: USA
ZIP: 02110-2804
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TELEFAX:
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                                     TELEPHONE:
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    617/542-8906
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Choi, Hueng-Sik
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Pred. No. 1
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Вþ

390 TQEEYALLTA 399

1 TSSEYQILTA 10

Query Match Best Local :

Local

Similarity 6; Conserv

Conservative

54.48;

Score 31; DB 5; Pred. No. 1.3e+02; 1; Mismatches 3

Length 484;

Indels

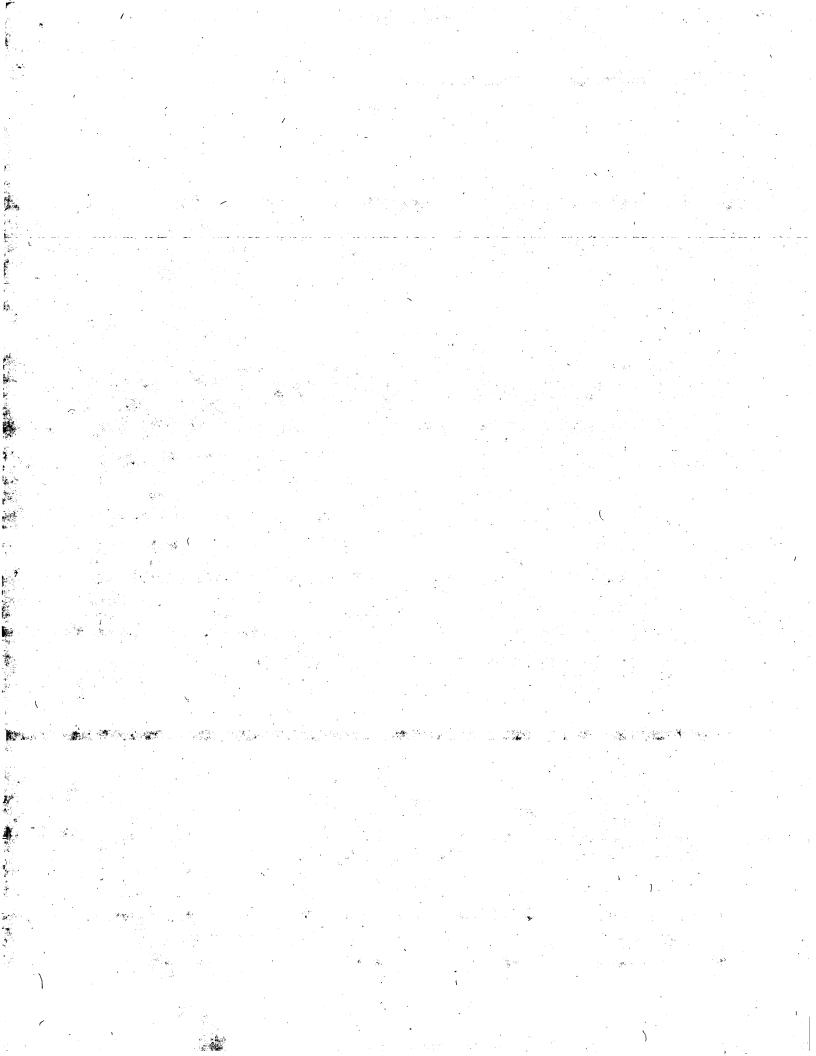
Gaps

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Query Match
Best Local Similarity
---hes 6; Conserv
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PCT-US95-16311-1
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                                                         TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 484 amin
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                                                                                                                                                                  FILING DATE: 13-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Clark Paul T:
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 00
            MOLECULE TYPE:
                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 617/542-5070
                                                                                                                                                                                                                                                                                                                                                                      ZIP: 02110-2804 COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
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                                                                                                               TELEPHONE: 617/542-8906
                                                                                                                                                                                                                              APPLICATION NUMBER: 08/3 FILING DATE: 13-JAN-1995
                            TOPOLOGY:
                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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225 Franklin Street, Suite 3100
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SYSTEM: PC-DOS/MS-DOS
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Search completed: June 4, 2001, 12:14:31 Job time: 224 sec



GenCore version Copyright (c) 1993 - 2000

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## ALIGNMENTS

RESULT A41357

Fc gamma (IgG) receptor I (high affinity)

form

b - human

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RESULT . 2
A39878
A39878
A39878
C gamma (IgG) receptor I-A (high affinity) precursor - human
N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Date: 30-Dec.1991 *sequence_revision 06-Sep-1996 *text_change 21-C;Caccession: A39878; I70304; B41357; S03018; I57525
R;van de Winkel, J.G.J.; Ernst. L.K.; Anderson, C.L.; Chiu, I.M.
J. Biol. Chem. 266, 13449-13455, 1991
A;Title: Gene organization of the human high affinity receptor for 1
A;Reference number: A39878; MUID:91302383
A;Molecule type: DNA
A;Residues: 1-374 <VAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 A;NOte: the authors translated the codon ACT for residue 25 as Ala C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein F;117-170/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000
C;Accession: A41357; S03019
C;Accession: A41357; S03019
C;Accession: A31357; S03019
C;Accession: A31357; B98
Science 243, 378-381, 1989
A;Title: Isolation and expression of functional high-affinity Fc receptor complementa
A;Reference number: A41357; MUID:89100284
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A; Residues: 1-344 < ALL2>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Allen, J.M.; Seed, B.
Nucleic Acids Res. 16, 11824, 1988
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A; Molecule type: mRNA
A; Residues: 1-344 <ALL1>
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Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  241 TSSEYQILTARR 252
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                                                                                                                      receptor for
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                                                                                                                   IgG, FcgammaRI (CD
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A;Molecule type: mRNA
A;Residues: 1-336 <RE
                                                                                                                           Science 260, 695-698,
                                                                                                                                                                                                                                                                                                                                 QY
                                                                                                                                                                                                                                                 RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Science 243, 378-381,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references:
                                                                                                                                             ;Prins,
                                                                                                                                                                        c gamma (IgG) receptor high affinity mouse; Species: Mus musculus (house mouse); Spate: 02-Jul-1996 #sequence_revision 02-Jul
                                                                                                         :Title: Linkage on chromosome 3 of
                                                                                                                                                           Accession: I48471
                                                       Status: preliminary; translated
                                                                           Accession:
                                                                                       Reference number: I48471; MUID:93242399
                                                                                                                                                                                                                                                                                                                                                                   Best Local
Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reference number:
Accession: I57525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Molecule type: mRNA
;Residues: 1-24,'S',26-337,'T',339-374 <AL2>;Cross-references: EMBL.X14356; NID:931331; |
;Perez, C.; Wietzerbin, J.; Benech, P.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Superfamily: Fc gamma receptor I; immunoglobulin homology 
Keywords: alternative splicing; glycoprotein; immunoglobu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Status: translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Molecule type: mRNA Residues: 1-24,'S',26-337,'T',339-374 ALL> Cross-references: GB:X14356; GB:M21091; NID:g31331;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title: Isolation and expression of functional high-affinity Reference number: A41357; MUID:89100284 Accession: B41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Allen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Title: Novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      117-170/Domain: immunoglobulin 293-313/Domain: transmembrane +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Residues: 1-8 <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Title: Nucleotide sequence of three CDNAs Reference number: S03018; MUID:89098339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cleic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Accession: S03018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references: GB:L03418; NID:g184840;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -15/Domain: signal sequence #status predicted 6-292/Domain: extracellular #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  3-313/Domain: transmembrane #status pj
78,152,159,163,195,240/Binding site:
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                                                                                                                                           J.B.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Invest.
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12; Conser
                                                                         I48471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wietzerbin, J.; Benech, P.D.
iol. 13, 2182-2192, 1993
cis-DNA elements involved in myeloid-cell-specific expression and gamma
                                                                                                                       Todd, J.A.;
495-698, 1993
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Seed, B.
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larity 100.0%;
Conservative
       EMBL: X70980;
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                                                                                                                                       Rodrigues, N.R.; Ghosh, S.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GB:M63835; NID:g180277; Doebele, R.; Pan, L.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OMIM: 146760
   NID: 9311748;
                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nomology <IMM2>
                                                       from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                              Score 57; -DB 1; 1
Pred. No. 0.00063;
; Mismatches 0;
                                                                                                       autoimmune
                                                                                                                                                                      predicted <TMM>
e: carbohydrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIDN: AAD13887.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIDN:CAA32537.1;
PIDN:CAA50311.1; PID:g311749
                                                                                                       diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                                                              Length 374
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                                                                                                                                      Hogarth, P.M.; Wicker, L.S.; Gaf:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Asn)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIDN:AAA35678.1; PID:g180279 Salmon, J.E.; Kimberly, R.P.
                                                                                                         and
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                                                                                                       defective Fc
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                                 A;Cross-references: GB:AE004638; GB:AE004091; NID:g9948116; PIDN:AAG05501.1; GSPDB:GNA;Experimental source: strain PAO1
                                                                   A; Molecule type:
A; Residues: 1-409
                                                                                                    A; Status: preliminary
                                                                                                                                    A;Title: Complete genome sequence of Pseudomonas aeruginosa A;Reference number: A82950; MUID:20437337
                                                                                                                                                                                                        adman,
                                                                                                                                                                                                                                                                   probable porin PA2113 [imported] - Pseudomonas aeruginosa (strain PAO1)
C;Species: Pseudomonas aeruginosa
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C;Keywords: immunoglobulin receptor
F;128-180/Domain: immunoglobulin hom
                                                                                                                     A;Accession: H83382
                                                                                                                                                                    Nature 406, 959-964,
                                                                                                                                                                                                                      R;Stover, C.K.; Pham, X.Q.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Keywords: immuno F; 127-179/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C; Species: Mus musculus (house mouse)
C; Date: 18-Jun-1993 #sequence_revision 18-Nov-1994
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A; Residues: 1-404 <SEA>
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Matches
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                                                                                                                                                                                      Olson, M.V
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8; Conser
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                                                                   <STO>
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                                                                                                                                                                                                    Brody,
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L.L.; (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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Pred. No. 15;
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Pred.
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r, S.N.; Folger,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <MMI>
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A;Introns: 21/1; 98/3; 130/3; 143/1; 178/3
C;Superfamily: Caenorhabditis elegans hypothetical protein C31H1.
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A;Title: Expression of the elcosapentaenoic acid
A;Reference number: 220764; MUID:97419510
A;Accession: T30185
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                                                                                                                                                                                                                                                                                                                                                                                            submitted to the EMBL Data Library, December 1995
A;Description: The sequence of C. elegans cosmid C31H1
A;Reference number: Z18395
                                   S46668
                                                 RESULT
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C;Species: Caenorhabditis elegans
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Cross-references: EMBL:U73935; NID:g2529413; PID:g2529420; PIDN:AAB81125.1
Experimental source: strain SCRC-2738
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                                                                                                                                                                               Query Match
Best Local
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    Caenorhabditis elegans

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A; Authors: Foulger, D.; FILLE, C., A; Hilbert, H.; Holsappel, S.; Hosono, S., L., lech, J.; Harwood, C.R.; Henaut, A,; Hilbert, H.; Kurita, K.; Lapidus, A.; Liech, J.; Karwood, C.R.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; Liu, H.; Masuda, S. Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Levine, A.; Liu, H.; Masuda, S. A; Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S. Y, M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; P., M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sekiguchi, J.; Sekowska, C.; Roche, R.; Scoffone, F.;                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Margot, P.; Mauel, C.; Karamata, D.
Mol. Microbiol. 12, 535-545, 1994
A;Title: The gene of the N-acetylglucosaminidase,
A;Reference number: S49389; MUID:95020588
                                                                                                                                                                                                                                                                                                           R;Kunst, F.; Ogasawara, N.; Moszer, I.; Albertini, A.M.; Alloni, G.; Azevedo, V.; Be
C.; Bron, S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.;
A.; Ehrlich, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari,
Nature 390, 249-256, 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Residues: 1-880 <MAR>
A; Cross-references: EMB
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C;Date: 15-Feb-1996 #sequence_revision 01-Mar-1996
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A; Residues: 1-433 < HUB>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 81-132; 1:
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                                 Tanaka,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.2.1.52)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NID: g1332633;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 35;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           October 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ed. No. 25;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disruption
                         Terpstra, P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-acetylglucosaminidase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lytD precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cosmid 9954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIDN: AAA18536.1; PID: g416160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PID:g1230646; MIPS:YDR277c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.
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                                                                                                                                                                                                                                                                                  S.; Galizzi, A.; Gal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (STD1) in
Yata,
                                                                                                                                         ; Portete
                                                                                                                                                                                                                                         Hullo,
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                                                                                                                                                                          S.; Mau
                                                                                                                                                                                                               Lardino
                                     Uchiya
                                                                                                   Scanl
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A;Cross-references: GB:AL009198; GB:AL123456; NID:g3242262; PIDN:CAA15760.1; A;Experimental source: strain H37Rv
                                                                                                  A;Authors: Sqares, R.; Súlston, J.E.; Taylor, K.; Whitehead, S.;
A;Title: Deciphering the biology of Mycobacterium tuberculosis f.
A;Reference number: A70500; MUID:98295987
A;Accession: F70972
                                                                                                                                                                                                                                                                                                                                                                                                                            밁
                                             A; Residues: 1-475 <COL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      닭
                                                                                                                                                                                   Nature 393,
                                                                                                                                                                                                                                                                                                                           probable amidase –
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references: EMBL:AL050352;
A;Experimental source: cultivar Col
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Keywords: glycosidase; hydrolase
F;1-27/Domain: signal sequence #status predicted
F;28-880/Product: beta-N-acetylhexosaminidase #st
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Introns: 27/2; 82/3; 168/3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Residues:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Status: preliminary; nucleic acid sequence not shown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Accession: F69654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Reference number:
                                                               ; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nypothetical
                                                                                                                                                                                                                  Connor, R.; Davies, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Hamris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Ho
                                                                                                                                                                                                                                                         ;Species: Mycobacterium tuberculosis;Date: 17-Jul-1998 #text_change 20-Jun-2000;Date: 17-Jul-1998 #sequence_revision, 17-Jul-1998 #text_change 20-Jun-2000;Accession: F70972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ubmitted to the Protein Sequence
                                                                                                                                                                               ijandream, M.A.; Rogers,
iture 393, 537-544, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ypothetical protein F27B13.170 - Arabidopsis thaliana
;Species: Arabidopsis thaliana (mouse-ear cress)
;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #te
                                                                              Status: preliminary; nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gene: ATSP:F27B13.170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Molecule type: DNA
Residues: 1-277 <BEV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reference number: Z16442
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cross-references: GB:Z99122; GB:AL009126; NID:g2636029; PIDN:CAB15595.1; PID:e1184484
Experimental source: strain 168
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257 TSTNYKILNASR 268
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 58. hes 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zimmermann, W.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           complete genome sequence of the Gram-positive bacterium Bacillus subtilis. number: A69580; \mathtt{MUID}:98044033
                                                                                                                                                                                                                                                                                                                     Mycobacterium tuberculosis (strain H37RV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      59.6%;
                                                                                                                                                                                                     J.; Rutter,
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63.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               52; GSPDB:GN00062; ATSP:F27B13.170 Columbia; BAC clone F27B13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 34;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score
Pred.
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No. 54;
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May 1999
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                                                                                                                                                                                                   s.,
                                                                                                                                          from the
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                                                                                                                                                            Barrell, B.G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                                                                                                                                                                 Squares,
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                                                                                                                                       complete
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                                                                                                                                                                                                                    Holroyd,
                    PID:g266165
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                                                                                                                                       genome
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В
                                   Qy
                                                                                                                                                                             C; Genetics: A; Gene: PA3
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C; Superfamily:
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                                                                            Matches
  123 SSVYQIRRARR 133
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A; Molecule type: DNA
A; Residues: 1-238 <STO>
A; Cross-references: GB: AEO04747; GB: AEO04091; NID: g9949362;
                                                                                                                                                                                                                                                                                              .; Lory, S.; Olson, M.V. Nature 406, 959-964, 2000
                                                                                                                                                        A; Experimental source: strain PAO1
                                                                                                                                                                                                                                            A;Reference number: A82950;
A;Accession: C83240
                                                                                                                                                                                                                                                                                                                                adman, S.; Yuan,
                                                                                                                                                                                                                                                                                                                                              R;Stover, C.K.; Pham, X.Q.; Erwin,
                                                                                                                                                                                                                                                                                                                                                                                              probable transcription regulator PA3249
C; Species: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            related to glucan 1,3-beta-glucosidase precursor [imported] - Neurospora crassa
N.Alternate names: protein B23L21.220
                                                                                                                                                                                                                                                                              A; Title: Complete genome
                                                                                                                                                                                                                                                                                                                                                                           C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Gene: NCSP: B23L21.220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Experimental source:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:AL356172;
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A; Residues: 1-863 <SCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Status: preliminary
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A; Accession: T49709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              submitted to the Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R; Schulte, U.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Accession: T49709
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                                                                                                                                                                                                                          ;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 6
                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 59.1
Best Local Similarity 77.1
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    228 SDDEYQYLTAKK 239
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2 SSEYQILTARR 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TSSEYQILTARR
                                                                                                                                                                                                                                                                                                         uan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Olson, M.V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aign, V.; Hoheisel, J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       indoleacetamide hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                            sequence of Pseudomonas aeruginosa PAO1, an opportunistic 50; MUID:20437337
                                                   57.9%;
72.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence Database, May
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     59.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          59.6%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .88;
                                  Score 33; DB Pred. No. 35; 0; Mismatches
                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 34; DB
Pred. No. 85;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; GSPDB:GN00116; NCSP:B23L21.220
B23L21; strain OR74A
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0; Mismatches
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Pred. No.
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    Pseudomonas aeruginosa (strain

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C;Species: Streptomyces coelicolor
C;Sate: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 31-Jan-2000
C.Accoscio. #7503
                                                                                                                                                                                                    A; Note: DKFZp762M186.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Gene: trpC2; SCOEDB:SCE8.04c
C;Superfamily: indole-3-glycerol-phosphate synthase; trpC homology
                                                                                     Query Match 57.9
Best Local Similarity 70.0
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                               ;Bloecker, H.; Boecher, M.; Brandt, P.; Mewes, H.W.; Weil, B.; Wiemann, S. bbmitted to the Protein Sequence Database, June 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 57.9%;
Best Local Similarity 87.5%;
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                                                                                                                                                                                                                                                                                     Status: preliminary
Molecule type: mRNA
Residues: 1-271 <AAA>
                                                                                                                                                                                                                                                                                                                                                      Reference number: Z25143
Accession: T50620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pothetical protein DKF2p762M186.1 - human (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Saunders, D.; Harris, D.; Bentley, S.D.; Parkhill, J.; Barrell, B.G.; Rajandream, M.A. bmitted to the EMBL Data Library, March 1999 Reference number: Z21604 Accession: T36303
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Species: Homo sapiens (man)
Date: 21-Jul-2000 #sequence_revision 21-Jul-2000 #text_change 21-Jul-2000
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Cross-references: EMBL:AL035654; PIDN:CAB38582.1; GSPDB:GN00070; SCOEDB:SCE8.04c
Experimental source: strain A3(2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Status: preliminary; translated from GB/EMBL/DDBJ Molecule type: DNA
                                                                                                                                                                                                                                      Cross-references: EMBL:AL359590
Experimental source: adult melanoma (MeWo cell line); clone DKFZp762M186
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34 31 54.4 403 1 T23O_CAEEL Q09474 caenorhabdi 35 31 54.4 437 1 FLHE_PSEPU O52256 pseudomonas 36 31 54.4 444 1 STD1_YEAST Q92794 saccharomyc 37 31 54.4 464 1 HSLU_BACHD Q9ka27 bacillus ha 38 31 54.4 477 1 TRPC_HAEIN P46451 haemophilus 39 31 54.4 480 1 HEP2_RABIT P4776 oryctolagus 40 31 54.4 680 1 SYG_BOMMO Q9451 bombyx mori 41 31 54.4 680 1 SYG_BOMMO Q04451 bombyx mori 42 31 54.4 699 1 YA76_SCHPO Q9761 schizosacch 43 31 54.4 791 1 HOX1_HALRO P57454 buchnera ap 44 31 54.4 794 1 HMMR_MOUSE ALIGNMENTS		_		_			.~			1.1			
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RESULT 1
FCG1\_HUMAN STANDARD; PRT; 374 AA.

ID FCG1\_HUMAN STANDARD; PRT; 374 AA.

AC P12314; P12315;
DT 01-OCT-1989 (Rel. 12, Created)
DT 01-OCT-1989 (Rel. 12, Last sequence update)
DT 01-OCT-2000 (Rel. 40, Last annotation update)
DT 01-OCT-2010 (Rel. 40, Last annotation update)
DE HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR (FC-GAMMA DE RI) (FCRI) (IGG FC RECEPTOR I) (CD64).

GN FCGRIAOR FCGRI OR FCGCI OR IGFRI.

OCT-2000 (Rel. 40, Last OCT-2000 (Rel. 40, Last OCT-2000 (Rel. 40, Last SHAPENITY IMMUNOGLOBULT) (RECEI) (IG FC RECEPTO) (RECEI) (IG FC RECEPTO) (RECEI) (RECE	S; Rel. 12, Created) Rel. 12, Last sequence update) Rel. 40, Last annotation update) IMMUNOGLOBULIN GAMMA FC RECEPTOR I	3G FC RECEPTOR I) (CD64).	(Human).	Eutheria; Primates; Catarrhini; Hominidae;	NCBI_TaxID=9606;		98339;	Seed B.;	ce of three cDNAs for the human high	Acids Res.	incress inco.	SEQUENCE FROM N.A.			ssion of functional high-affinity FC	Science 243:378-381(1989)	CTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS	CEPTOR.	I - ALTERNATIVE DECOMINES: O TROPODEMS: A CHOWN HERE) AND	PRODUCED BY ALTERNATIVE SPLICING.	- TISSUE SPECIFICITY: N	I- SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2-TYPE	WWW="http://www.ncbi.nlm.nih.gov/prow/cd/cd64.htm".		uced throug	and the	non-profit institutions as long as its cont	ified and this statement is not removed.	tities requires a license agreement (See	nd an email to license@isb-	W143EC (23033E37 1	CAA3233/.1;	X14355; CAA32536.1;	; X14356; CAA32536.1; ; X14355; CAA32536.1; S03018; S03018.	X14355; CAA32536.1; X14355; CAA32536.1; X03018; X03018. X03019; X03019. X41357; X41357	;; X14355; CAA32536.1; ;; X14355; CAA32536.1; ; S03018; S03018. S03019; S03019. S03019; S03019. A41357; A41357.	, A14355; CAA32536.1; ; X14355; CAA32536.1; S03018; S03018. S03019; S03019. A41357; A41357. B41357; B41357. B41357; B41357.	X44355; CAA3236.1; X14355; CAA32536.1; S03018; S03019. S03019; S03019. A41357; A41357. B41357; B41357. B41357; B41357. P12319; IALT.	, X14355 (CAX32536.1; ; X14355; CAX32536.1; \$03018; \$03019. \$03019; \$03019. \$03019; \$03019. \$41357; \$41357. \$41357; \$41357. \$41357; \$41357. \$146760; \$18003006; \$1600047.
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Best Local Similarity 100 Matches 12; Conservative
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Sears D.W., Osman N., 7
"Molecular cloning and
receptor for IgG.";
                                                             "Structure
                                                                               MEDLINE=92166399;
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Mus musculus (Mouse).

Chordata;
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15-JUL-1999 (Rel.
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                                          mman N., Kozak C.A., McKenzie I.F., Hogarth P.M., 
ttructure and mapping of the gene encoding mouse 
mma RI and chromosomal location of the human Fc 
Immunol 148:1570-1575(1992).
                                                                                                        Immuno1.
                                                                                                                                                                                                                                                                                                                    1 TSSEYQILTARR
 TISSUE SPECIFICITY: N
SIMILARITY: CONTAINS
          AFFINITY RECEPTOR.
SUBCELLULAR LOCATION: TYPE I MEMBRANE
TISSUE SPECIFICITY: MACROPHAGE-SPECICI
                                  FUNCTION: BINDS TO THE FC
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Rel. 22, Last sequence update)

Rel. 38, Last annotation update)

IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR (FC-GAMMA
GG FC RECEPTOR I).
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                                                                                                                                         PubMed=2136886;
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MTH1_YEAST
P35198;
01-FEB-1994
01-FEB-1994
01-OCT-1996
           Johnston M., Andrews S., I
Fayello A., Fulton L., Gat
Kucaba T., Hallsworth K.,
                                                                                         MEDLINE-94158870; PubMed-8114728;
Hubbard E.J.A., Jiang R., Carlson M.
"Dosage-dependent modulation of gluc
Saccharomyces cerevisiae.";
Mol. Cell. Biol. 14:1972-1978(1994);
                                                                                                                                                                                                                                                     MTH1 PROTEIN.
MTH1 OR YDR277C OR D9954.12.
                                     STRAIN=S288C / AB972;
Johnston M., Andrews S.,
                                                                 SEQUENCE FROM N.A
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                                                                                                                                                                                                    Saccharomycetales;
NCBI_TaxID=4932;
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PIR; A43511; A43511.
PIR; A46480; A46480.
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., Brinkman R., Cooper J., Ding H., Gattung S., Greco T., Kirsten J., K., Hawkins J., Hillier L., Jier M., Langston Y., Latreille P., Le T.,
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IG-LIKE C2-TYPE DOMAIN.

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IG-LIKE C2-TYPE DOMAIN.
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Rashid M.H., Mori M., Sekiguchi J.;
"Glucosaminidase of Bacillus subtilis: cloning, regulation, primary structure and biochemical characterization.";
Microbiology 141:2391-2404 (1995).

HICROBION: CELL WALL HAYDROLASE NOT INVOLVED IN CELL AUTOLYSIS. IT HYDROLYZES THE BETA-1,4 GLYCAN BOND BETWEEN THE N-ACETYLGUCOSAMINYL AND THE N-ACETYLMURAMOYL RESIDUES IN THE GLYCAN
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01-FEB-1995 (Rel. 31, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
BETA-N-ACETYLGLUCOSAMINIDASE PRECURSOR (EC 3.2.1.-).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Margot P.,
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SUBUNIT: HOMODIMER.
SUBCELLULAR LOCATION:
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gene of the N-acetylglucosaminidase,
wall hydrolase not involved in vegeta
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S0002685; MTH1.
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                       http://www.isb-sib.ch/announce/
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Best Local S
Matches 7
                                                                                                                                                                                               Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy Oliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J. Rutter S., Seeger K., Skelton S., Squares S., Sqares R., Sul Taylor K., Whitehead S., Barrell B.G.; "Deciphering the biology of Mycobacterium tuberculosis from complete genome sequence."; Nature 393:537-544(1998).

-i- CATALYTIC ACTIVITY: A MONOCARBOXYLIC ACID AMIDE + H(2)0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AMI4_MYCTU
050404;
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Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C.
Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia
Badcock K., Basham D., Brown D., Chillingworth T., Connor
Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N.,
                                                                                                   the European Bioinformatics Institute. The suse by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
PUTATIVE AMIDASE AMID (EC 3.5.1.4).
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EMBL; D45048; BAA08089.1;
                                          TubercuList; Rv3375;
                                                         EMBL; AL009198; CAA15760.1;
                                                                              or send
                                                                                            entities requires a license
                                                                                                                                      between the Swiss Institute of Bioinformatics
                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                            Actinomycetales;
NCBI_TaxID=1773;
                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Firmicutes;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REPEAT
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SIMILARITY: BELONGS TO THE AMIDASE FAMILY.
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BETA-N-ACETYLGLUCOSAMINIDASE.
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Z_ENTHI STANDARD; PRT; 288 AA. 03077; 01-JUL-1993 (Rel. 26, Created) 01-JUL-1993 (Rel. 26, Last sequence update) 01-NOV-1995 (Rel. 32, Last annotation update)
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Q9Z4X0;
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                             modified and this statement is not removentities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                              Tryptophan biosynthesis; Lyase; Decarboxylase; Antibiotic biosynthesis.
                                                                                                                                                                                                                                                                                                           EMBL; AL035654; CAB38582.1; HSSP; P00909; 1PII.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rajandream M.A.;
Submitted (MAR-1999) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION: THE FUNCTION OF THE SECOND TRP OPERON IN S.COELICOLOR TO PRODUCE TRP FOR THE BIOSYNTHESIS OF CALCIUM-DEPENDENT ANTIBIOTIC (CDA).
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                                                                                                                                                                                                                                                                                                                                                                                  the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacteria; Firmicutes; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
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30-MAY-2000 (Rel. 39, Last annotation update)
INDOLE-3-GLYCEROL PHOSPHATE SYNTHASE 2 (EC 4.1.1.48) (IGPS)
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SIMILARITY: BELONGS TO THE TRPC FAMILY.
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77
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Pred. No. 14;
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EMBL; L06065; AAA16054.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                use by non-profit institutions as long modified and this statement is not removed.
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MCDILNE-94043255; PubMed-8226970;
MCCOY J.J., Mann B.J., Vedvick T.S.,
Petri W.A. Jr.;
                                                                                                                                                                                                                                                                     PIR; A48452;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Molecular cloning of cDNA and genomic sequences coding for the kilodalton subunit of the galactose-inhibitable lectin of pathos Entamoeba histolytica.";
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J. Biol. Chem. 268:24223-24231(1993)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBUNIT:
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                     GALACTOSE INHIBITABLE LECTIN 35
SUBUNIT.
S -> A (IN REF. 2).
E -> K (IN REF. 2).
E -> G (IN REF. 2).
DECF366D3DAFA24B
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                                 Score 33;
Pred. No.
                           Mismatches
                                 16;
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RESULT
PARB_C!
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005190;
01=0CT-2000
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MEDLINE=97207014; PubMed=9054507; Mohl D.A., Gober J.W.;
                           SEQUENCE FROM N.A STRAIN=NA1000;
                                                                    NCBI_TaxID=76;
                                                                                  Caulobacter
                                                                                              Bacteria; Proteobacteria;
                                                                                                            Caulobacter crescentus.
                                                                                                                                         CHROMOSOME
                                                                                                                                                     01-OCT-2000
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(Rel. 40, Last ann
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                                                                                            alpha subdivision; Caulobacter group;
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cycle-dependent polar localization of chromosome partitioning

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Pfam; PF02195; ParBc; 1.
Chromosome partition; DNA-binding.
Chromosome partition; DNA-binding.
293 AA; 31910 MW; E7D557D90F61469C CRC64;
                                                                                            This SWISS-PROT entry is copyright: It is produced through a between the Swiss Institute of Bioinformatics and the EMBI the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and f
                                                                                                                                                                                                                                                                                                     "Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. I. Sequence features in the 1 Mb region from map positions 64% to 92% of the genome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
PUTATIVE METHIONINE AMINOPEPTIDASE C (EC 3.4.11.18) (MAP) (PEPTIDASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SYNY3
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                                                                                                                                                                                                                                           -I- FUNCTION: REMOVES THE AMINO-TERMINAL MI
PROTEINS (BY SIMILARITY).
-I- CATALYTIC ACTIVITY: L-METHIONYLPEPTIDE
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P53581;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proteins in Caulobacter crescentus."; Cell 88:675-684(1997).
                         EMBL; D64003; BAA10478.1; -. HSSP; P07906; IMAT.
                                                                                                                                                                                                                                                                                                                                                Sugiura M., Tabata S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Synechocystis sp. (strain PCC 6803).
Bacteria; Cyanobacteria; Chroococcales; Synechocystis
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                                                                      or send
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                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SLL0555
                                                                                                                                                                                                                                                                                                                                                            Kaneko T., Tanaka A., Sato S., Kotani H.,
                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-96127529; PubMed-8590279;
                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=1148;
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SIMILARITY: BELONGS TO PEPTIDASE FAMILY M24A;
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                                                                                                                                                                                        MAP FAMILY 1.
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6; Conserv
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an email to license@isb-sib.ch).
                                                                                     requires a license agreement
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ARITY: BELONGS TO THE PARB PROTEIN FAMILY.
                                                                    email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Best Local
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METAL 147 147
METAL 158 158
METAL 221 221
                                                                                                                                                                   use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and foentities requires a license agreement (See http://www.isb-slb.or send an email to license@isb-slb.ch).
                                                                                                                                                                                                                                                                                                              "Cloning and tissue distribution of the human G protein beta 5 cDNA Biochim. Biophys. Acta 1402:288-291(1998).

-- FUNCTION: GRANINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE INVOLVED AS A MODULATOR OR TRANSDUCER IN VARIOUS TRANSMEMBRANE SIGNALING SYSTEMS. THE BETA AND GAMMA CHAINS ARE REQUIRED FOR T GTPASE ACTIVITY, FOR REPLACEMENT OF GDP BY GTP, AND FOR G PROTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHAIN 5).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-1998 (Rel. 36, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               014775;
15-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HUMAN
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                     PROSITE; PS00678; WD_REPEATS_1; 3. PROSITE; PS50082; WD_REPEATS_2; 6. PROSITE; PS50294; WD_REPEATS_REGION;
                                                                                                                                 EMBL; AF017656; AAC63826.1; MIM; 604447; -.
                                                                                                                                                                                                     the European Bioinformatics Institute. The use by non-profit institutions as long
                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
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InterPro; IPR002467;
                                                        PRINTS; PR00319; GPROTEINB.
PRINTS; PR00320; GPROTEINBRPT
                                                                                                                                                                                                                                                                MEDLINE=98269832; PubMed=9606987
                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Transducer;
REPEAT
                                                                                   InterPro; IPR001680;
Pfam; PF00400; WD40;
                                                                                                        InterPro; IPR001632;
                                                                                                                      HSSP; P04901; 1GP2
                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          METAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  292 TATGYELLTDRR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 TSSEYQILTARR 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HUMAN
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Repeat;
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                                                                                                                                                                                                                                                                                                                                                                                         Lombardi S.J.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STANDARD;
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repeat; Multigene
WD 1.
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                                                                                                                                                                                                                                                                                                                                                                                         Cockett M.I.;
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Pred. No. 17;
4; Mismatches
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COBALT
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BBALT (BY SIMILARITY).

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17;
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            family
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Best Local Similarity
Watches 7; Conserv
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                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                              Betty M., Harnish S.W., Rhodes K.J., Cocke "Distribution of heterotrimeric G-protein the rat brain.";
                                                use by non-profit institutions as long a
modified and this statement is not removed.
entities requires a license agreement (See )
                                                                                                                                        -!- SUBUNIT: G PROTEINS ARE COMPOSED OF 3 UNITS (ALPHA, E-!- TISSUE SPECIFICITY: BRAIN AND AT MUCH REDUCED LEVELS -!- SIMILARITY: CONTAINS 7 WD REPEATS (TRP-ASP DOMAINS).
                                                                                                                                                                                   Neuroscience 85:475-486(1998).

-i- FUNCTION: GUANINE NUCLEOTIDE-BINDING PROTEINS (G PROTEINS) ARE INVOLVED AS A MODULATOR OF TRANSDUCER IN VARIOUS TRANSMEMBRANE SIGNALING SYSTEMS. THE BETA AND GAMMA CHAINS ARE REQUIRED FOR THE GTPASE ACTIVITY, FOR REPLACEMENT OF GDP BY GTP, AND FOR G PROTEIN STEFECTOR INTERACTION.
                                                                                                                                                                                                                                                                                                                                                                           SPECIES-Rat; STRAIN-SPRAGUE-DAWLEY; TISSUE-Brain Levay K., Slepak V.Z.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Watson A.J., Katz A., Simon M.I.,
"A fifth member of the mammalian G-protein beta-subunit family.
Expression in brain and activation of the beta 2 isotype of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GBB5_MOUSE STANDARD; PRT; 353 AA.
P54314; 035354;
01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
GUANINE NUCLEOTIDE-BINDING PROTEIN BETA SUBUNIT 5 (TRANSDUCIN BETA
                                      or send
                                                                                                                                                                                                                                                                                                             SPECIES=Rat; STRAIN-SPRAGUE-DAWLEY; MEDLINE-98283588; PubMed-9622245;
                                                                                                                                                                                                                                                                                                                                                                 Submitted (MAY-1997) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mus musculus (Mouse), and Rattus norvegicus (Rat). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I Eukaryota; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REPEAT
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                                  s requires a license agreement (s an email to license@isb-sib.ch)
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70.0%;
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Pred.
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                                                                                                                                                                                                                                                                                             Cockett
                                             (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                                                                                beta
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ta and
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                                                                                                                                                                                                                                                                               gamma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murinae;
                                                                                                                                                                      BETA & GAMMA)
                                                                                                                                                        IN KIDNEY
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EMBL; EMBL;

L34290; AZ AF001953;

AAA93084

AAB59974.1;

Qy

1 TSSEYQILTA

Matches Query Match Best Local:

Local Similarity hes 7; Conser

Conservative

0;

Mismatches

Indels

0;

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0;

57.9%;

Score 33; DE Pred. No. 24;

DB 1;

Length 413

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Best Local S
Matches 7
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15-JUL-1999 (Rel. 3
15-JUL-1999 (Rel. 3
HYPOTHETICAL 49.0 K
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045435;
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REPEAT
SEQUENCE
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REPEAT
REPEAT
                                                                                       use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                          15-JUL-1999
15-JUL-1999
15-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                LT 12
_CAEEL
  SEQUENCE
           Pfam; PF01062; Worm_family_8; Hypothetical protein.
                                           EMBL; Z81074; CAB03043.1; WormPep; F32B6.9; CE09864.
                                                                            entities requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
                                                                                                                                                                   Submitted (OCT-1996) to the EMBL/GenBank/DDBJ databases -!- SIMILARITY: BELONGS TO THE BESTROPHIN FAMILY.
                                                                                                                                                                                                      SEQUENCE FROM N.A. STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Nemato
Rhabditidae; Peloderinae;
                                  InterPro; IPR000615;
                                                                                                                                                                                            Basham V.;
                                                                                                                                                                                                                                         NCBI_TaxID=6239;
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REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE; PS00678; WD_REPEATS_1; 3. PROSITE; PS50082; WD_REPEATS_2; 6. PROSITE; PS50294; WD_REPEATS_REGION;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR001632; -
InterPro; IPR001680; -
Pfam; PF00400; WD40; 7
                                                                                                                                                                                                                                                                         Caenorhabditis elegans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REPEAT
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HSSP; P04901; 1GP2
MGD; MGI:101848; Gr
                                                                                                                                                                                                                                                                                                                                                                                                            162 TNSDMQILTA 171
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61
103
153
153
193
237
281
281
323
323
3A3;
11 protein.
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                                                                                                                                                                                                                                                                                          38, Created)
38, Last sequence update)
38, Last annotation update)
KDA PROTEIN F32B6.9 IN CHROMOSOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  :; WD repeat; 1
91 WD
133 WD
181 WD
225 WD
267 WD
311 WD
4.8965 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38731 MW;
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70.0%;
                                                                                                                                                                                                                                                 Caenorhabditis
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Pred. No. 20;
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 A66E69A83C78790B CRC64;
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165

TEKEYQILEA 174

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TRPG_YEAST
ID TRPG_YEAST
AC P00937;
   RROCCOSEE
                                                                                                                                                                                                                                                                                                                                                                                                                                               Aebi M., Furter R., Prantl F., Niederberger P., Huetter R.; "Structure and function of the TRP3 gene of Saccharomyces cerevisiae: analysis of transcription, promoter sequence, and sequence coding for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-84162082; PubMed-6323449;
Zalkin H., Paluh J.L., van Cleemput M., Moye W.S.
"Nucleotide sequence of Saccharomyces cerevisiae
encoding bifunctional anthranilate synthase: indephosphate synthase.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saccharomyces cerevisiae (Baker's yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                               entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                              the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed
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                                                                                                                                                                                                                                                                                                                                                                                                                                   a glutamine amidotransferase.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tzermia M., Horaitis O., Alexandraki D.;
"The complete sequencing of a 24.6 kb segment of yeast chromosome identified the known loci URA1, SAC1 and TRP3, and revealed 6 new open reading frames including homologues to the threonine dehydratases, membrane transporters, hydantoinases and the phosphoilpase A2-activating protein.";
yeast 10:663-679(1994),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. Pohl T.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TRP3 OR YKL211C
                                                                                                                                                  This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN-S288C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Saccharomycetales;
NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AMIDOTRANSFERASE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-1999 (Rel. 38,
ANTHRANILATE SYNTHASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 1-280 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted
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5-JUL-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biol.
                                                                                                                                                                                                                                                                              CATALYTIC ACTIVITY: 1-(2-CARBOXYPHENYLAMINO)-1-DEOXY-D-RIBULOSE 5-PHOSPHATE = 1-(INDOL-3-YL)GLYCEROL 3-PHOSPHATE + CO(2) + H(2)C PATHWAY: FIRST AND FOURTH STEDE IN BIOSYNTHESIS OF TRYPTOPHAN. SUBUNIT: TETRAMER OF TWO COMPONERYS I AND TWO COMPONENTS II. INDUCTION: BY TRYPTOPHAN STARVATION.

MISCELLANEOUS: COMPONENT I CATALYZES THE FORMATION OF ANTHRANILAMICCELLANEOUS: COMPONENT II.
                                                                                                                                                                                                MISCELLANEOUS: YEAST COMPONENT II CARBOXYL END ALSO INDOLE-3-GLYCEROL PHOSPHATE SYNTHASE ACTIVITY. SIMILARITY: THE GATASE DOMAIN BELONGS TO TYPE-1 GLUT
                                                                                                                                                                                                                                                 USING AMMONIA RATHER THAN GLUTAMINE, WHEREAS GLUTAMINE AMIDOTRANSFERASE ACTIVITY.
                                                                                                                                                                                                                                                                                                                                                                                     CATALYTIC ACTIVITY: CHORISMATE + L-GLUTAMINE PYRUVATE + L-GLUTAMATE.
                                                                                                                                                                                  AMIDOTRANSFERASES
                                                                                                                            SWISS-PROT entry is copyright. It is producen the Swiss Institute of Bioinformatics
K01386;
X75951;
                                                                                                                                                                                                                                                                                                                                                                                                                     Genet. 8:165-172(1984)
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Last annotation update)
COMPONENT II (EC 4.1.3.27)
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                                                           (See http://www.isb-sib.ch/announce/
                                                                                              There are no rest
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Best Local
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EMBL; M36300; AAA34450
PIR; S38049; NNBY2.
PIR; S44327; S44327.
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Q05754;
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PRINTS; PRO0097; ANTSNTHASEII.

PROSITE; PSO00442; GATASE_TYPE_I; 1.

PROSITE; PS00614; IGPS; 1.

Tryptophan blosynthesis; Transferas
Glutamine amidotransferase.
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InterPro;
                                                                                                                           McMahon A., Geertman R., Sabban E.L.;
McMahon A., Geertman R., Sabban E.L.;
"Rat dopamine beta-hydroxylase: molecular cloning a
"Rat dopamine beta-hydroxylase: molecular cloning a
                                                                                                                                                                                                                                                           01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation updat
                                                -1- COFACTOR: PQQ, COPPER, AND ASCORBATI
-1- PATHWAY: CATECHOLAMINE BIOSYNTHESIS.
-1- SUBUNIT: HOMOTETRAMER.
                                                                            SEQUENCE FROM N.A. MEDLINE-90219125;
                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                      Rattus norvegicus (Rat).
                                                                                                                                                                                                                                 DBH
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DOPAMINE BETA-MONOOXYGENASE PRECURSOR (EC 1.14.17.1) (DOPAMINE BETA-
                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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        SUBCELLULAR LOCATION: EXISTS AS BOTH A SOLUBLE AND MEMBRANE-BO FORM IN CHROMAFFIN GRANULES (THE MEMBRANE-BOUND FORM MAY BE ANCHORED BY AN UNCLEAVED SIGNAL SEQUENCE).

TISSUE SPECIFICITY: CHROMAFFIN GRANULES OF THE ADRENAL MEDULLA
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M36300;
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IPR002385;
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AAA34450.1;
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Rodentia;
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77.8%;
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                                                                    AND ASCORBATE.

    Mismatches

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Pred. No.
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Sciurognathi; Muridae;
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SYMPATHETIC NERVOUS
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e mRNA
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This

MONOOXYGENASE FAMILY

SIMILARITY: BELONGS TO THE COPPER TYPE II, ASCORBATE-DEPENDENT

entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).

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the European Bioinformatics Institute.

SWISS-PROT entry is copyright. It is produced through a collaboration sen the Swiss Institute of Bioinformatics and the EMBL outstation .

There are no rest

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RESULT 15
CJBA_BACUH
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Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                                                           CUBA_BACUH STANDARD; PKI; UU2.....
086170; 086170; 01-OCT-2000 (Rel. 40, Created)
01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
01-OCT-2000 (Rel. 40, Last annotation update)
PESTICIDIAL CRYSTAL PROTEIN CRY19BA (INSECTICIDAL DELTA-ENDOTOXIN)
PESTICIDIAL CRYSTALINE ENTOMOCIDAL PROTOXIN) (78 KDA CRYSTAL
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CARBOHYD
SEQUENCE
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MEDLINB=98369730; PubMed=9704107;
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MEDLINB=98369730; PubMed=9704107;
MEDLINB=98369730; PubMed=9704107;
MEDLINB=98369730; Pu
                                                                                                                                                                                                                                              Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillus/Staphylococcus group; Bacillus: NCBI_TaxID=132266;
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CRY19BA OR CRYXIXB(A).
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PROSITE; PS00084; CU2_MONOOXYGENASE_1;

PROSITE; PS00085; CU2_MONOOXYGENASE_2;
                                                                                                                                                                                                                                                                                                                              Bacillus thuringiensis
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BRITHBLIAL CELLS OF MOSQUITOS.

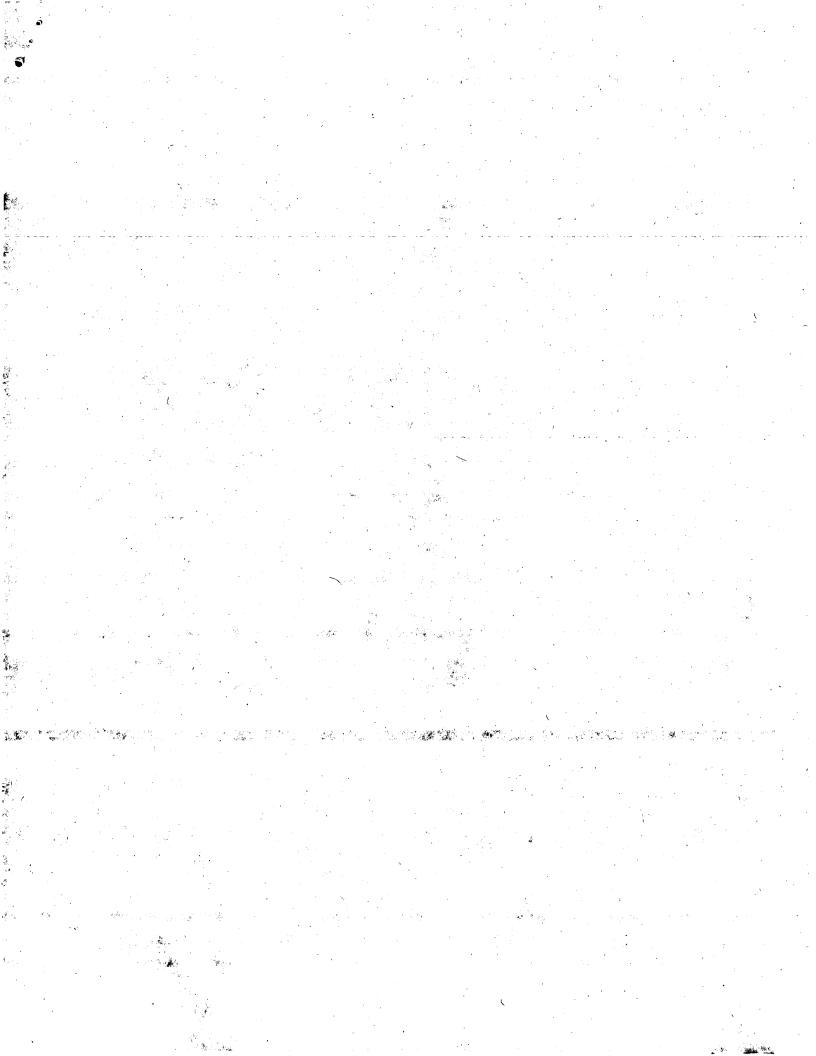
DEVELOPMENTAL STAGE: THE CRYSTAL PROTEIN IS PRODUCED DURING SPORULATION AND IS ACCUMULATED BOTH AS AN INCLUSION AND AS OF THE SPORE COAT.
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InterPro; IPR001178; -. Pfam; PF00555; endotoxin; 1. Toxin; Sporulation; Plasmid. SEQUENCE 682 AA; 78490 MW;
                                                                                                                                                                                                                  use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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## ALIGNMENTS

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RESULT
Q92495
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Best Local Similarity 100.0%;
Matches 12; Conservative 0;
Q92495
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J. Clin. Invest. 90:2102-2109(1992).
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Porges A.J., Redecha P.B., Doebele R.,
Kimberly R.P.;
                                                                                                                                                                                                                            INTERPRO; IPRO03006; -. PFAM; PF00047; ig; 3. PRODOM; PD002534; -; 1. SEQUENCE 374 AA; 4263
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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HSSP; P12319; 1ALT.
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SEQUENCE FAVO.

STRAIN-CV. COLUMBIA;

STRAIN-CV. COLUMBIA;

Lee J.M., Vaysberg M., Sakano H., Lenz C., Liu S.X., Pham P.,

Lee J.M., Vaysberg M., Sakano H., Lenz C., Liu S.X., Pham P.,

Toriumi M., Yu G., Chin C., Chiou J., Choi E., Chung M., Gonzalez

Toriumi M., Yu G., Chin C., Chiou J., Choi E., Chung M., Gonzalez

Howng B., Liu A., Altafi H., Brooks S., Buehler E., Chao Q., Conn

Connau A B., Hansen N.F., Johnson-Hopson C., Khan S., Kim C., Lam
                                                                                                                                                                                                                                                    Eukaryota; Viridiplantae; Embryophyta; Magnoliophyta; eudicotyledons; core eud Brassicales; Brassicaceae.
                                                                                                                                                                                                                                                                                                                                                                                                                        Q9LPHO;
01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q9LPH0
                                                                                                                                                                                                                                      NCBI_TaxID=3702;
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01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)
FC GAMMA RECEPTOR TYPE I (FC GAMMA RIB-FC GAMMA RECEPTOR).
CD64 OR FC-GAMMA>RIB.
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01-FEB-1997
01-FEB-1997
01-OCT-2000
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EMBL; M91555; AAA58414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Definition of interferon gamma receptor gene (Fc ga
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Mammalia; Eutheria;
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ilarity 100.
Conservative
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AAA58414.1; JOINED:
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Last sequence up
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dicots; Rosidae;
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Matches 7
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01-OCT-2000
01-OCT-2000
01-OCT-2000
    SEQUENCE FROM N.A.
Sherburne C.K., Lawley T.D., Gilmour M.W., Blat Sherburne C.K., Lawley T.D., Gilmour M.W., Blat Grotbeck E., Rose D.J., Taylor D.E.,
"The complete DNA sequence and analysis of R27, from Salmonella typhi that is temperature sensi Nucleic Acids Res. 28:217-2186(2000).
EMBL, AF250878; AAF69908.1;
                                                                                   "The sequence of BAC T3F20 from Arabidopsis the Submitted (JUN-2000) to the EMBL; AC018748; AAF78435.1; SEQUENCE 390 AA; 45480 vm
                                                                                                                   Salmonella
                                                                              NCBI_TaxID=601;
                                                                                                                             R0070
                                                                                                                                    HYPOTHETICAL
                                                                                                                                                                                  Q9L5R1
                                                                                                                                                                                                                                                                                                                            "Molecular cloning and identification of full-length cDNA high affinity Fc receptor for bovine IgG (Fc gamma RI)."; Vet. Immunol. Immunopathol. 75:151-159(2000).
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01-OCT-2000 (TrEMBLrel.
FC GAMMA RECEPTOR I.
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
Bovidae; Bovinae; Bos.
NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                Bos taurus (Bovine).
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01-OCT-2000
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PFAM; PF01377; Thioester_dehyd; 2.
PROSITE; PS00606; B_KETOACYL_SYNTHASE; UNKNOWN_1.
PROSITE; PS00881; PROTEIN_SPLICING; UNKNOWN_1.
PRODOM; PD013327; -; 2.
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STRAIN-SCRC-2738;
MEDLINE-97419510; PubMed-9274025;
Takeyama H., Takeda D., Yazawa K., Y
Takeyama hour the elcosapentaenoic
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                                                                                                                                                         Drosophila melanogaster
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EMBL; U73935; AABB1125.1;
HSSP; P18391; 1MKA.
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SEQUENCE FROM N.A. STRAIN-BERKELEY;
                                                                   NCBI_TaxID=7227;
                                                                                          Ephydroidea;
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                                                                                     ster (Fruit fly).
Arthropoda; Tracheata; Hexapoda; Insecta;
; Endopterygota; Diptera; Brachycera; Muschilidae; Drosophila.
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ine cyanobacterium, Synechococcus
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A Spier E., Spradling A.C., Stapleton M., Strong R., Sun E.,

Svirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,

A Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

A Williams S.M., Woodage T., Weinstock G.M., Yang S., Yao Q.A.,

A Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,

A Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu X., Smith H.O.

A Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.;

"The genome sequence of Drosophila melanogaster.";

Science 287:2185-2195(2000).

R EMBL; ABC003514; ABF49045.1; -

R FLYBASE; FBgn0036960; CG13814.

Q SEQUENCE 104 AA; 11491 MW; 2620F15C7CBFB431 CRC64;
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Q18337;
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01-NOV-1996 (7)
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COSMID C31H1.
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                                                                                                                                 Caenorhabditis elegans.
Eukaryota; Metazoa; Nemato
Rhabditidae; Peloderinae;
NCBI_TaxID=6239;
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Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A.
George R.A., Lewis S.E., Richards S., Ashburner M.,
Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., (
Wilson R., Ainscough R., Anderson K., Bayn
Bonfield J., Burton J., Connell M., Copsey
Craxton M., Dear S., Du Z., Durbin R., Fav
Gardner A., Green P., Hawkins T., Hillier
                                                                         MEDLINE=94150718; PubMed=7906398;
                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                     C31H1.2
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                                                                                                                                                                                Nematoda;
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                                                                                                                                                         oda; Chromadorea; Rhabditida; Rhabditoidea; Caenorhabditis.
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                                                                                                                                                                                                                                                                                                                                        PRT;
                                  Copsey T., Cooper J.,
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                  Favello A., Fulton
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8. 9
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                                                                                                                                                                                                                                                          update:
  Jier M., Johns
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X., Smith H.O.,
                                         Coulson A.,
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Query Match
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                                                                                            Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Durbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W., Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M., Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J., Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C.,
                                                                                                                                                                                                                                                       Wan K. H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G., Abril J.F., Agbayani A., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Borkova D., Botchan M.R., Bouck J., Bokstein P., Brottler P., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P., Chandra I., Deel C., Davenport L.B., Davies P., Chandra A., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng Z., Mays A.D., Dew I., Dietz S.M., Davies P., Chandra R., Deng R., Chandra R., Chandra R., Chandra R., Chandra R., Chandra R., Ch
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01-MAY-2000
01-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Eukaryota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha;
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q9W458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=20196006;
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                       Lasko P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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Lighthing J., Lloyd C., Mcmurray A., Mortimore B., O'Call
Parsons J., Percy C., Rifken L., Roopra A., Saunders D.,
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J
Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Waters
Watson A., Weinstock L., Wilkinson-Sproat J., Wohldman P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "2.2 Mb of contiguous nucleotide sequence from chromosome III of c.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (JAN-1996) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fe
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           M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A., B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 (TremBLrel. 13, 0 (TremBLrel. 13, 0 (TremBLrel. 14,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=10731132;
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58.3%;
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McLeod M.P.
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No.
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RESULT 11 Q9SZR4 ID Q9SZR4

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Best Local
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Q9WU16;
Q1-NOV-1999 (TrEMBLrel. 12, Created)
Q1-NOV-1999 (TrEMBLrel. 12, Last sequence up)
Q1-CCT-2000 (TrEMBLrel. 15, Last annotation)
Q1-CCT-2000 (TremBLrel. 15, Last annotation)
                                                                                                                                          NON_TER
                                                                                                                                                                  PFAM; PF00243; NGF;
                                                                                                                                                                                EMBL; AF104240; AAD21011.1;
HSSP; P20783; 1B8K.
INTERPRO; IPR002072; -.
                                                                                                                                                                                                                          Jansen H.T., Lehman M.N., Stevens P.J.;
"Golden Hamster Neurotrophin and Neurotrophin
Submitted (NOV-1998) to the EMBL/GenBank/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J., Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A., Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L., Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Zhu X., Smith H.O., Zibbs R.A., Myers E.W., Rubin G.M., Venter J.C.; "The genome sequence of Drosophila melanogaster."; Science 287:2185-2195(2000)
                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
Jansen H.T., Lehman M.N.,
                                                                                                                                                                                                                                                                                              NCBI_TaxID=10036;
                                                                                                                                                                                                                                                                                                                           Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                     Mesocricetus auratus (Golden hamster)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INTERPRO; IPR000822; -. PFAM; PF00096; zf-C2H2; 2. PFAM; PF00651; BTB; 1.
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                                                                                                                                                                                                                                                                                                             Mesocricetus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; AE003435; AAF46100.1; -. FLYBASE; FBgn0029822; CG12236.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Merkulov G., Milshina N.V., Mount S.M., Moy M., Murphy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               363 SSEYEILT 370
 41
                       1 TSSEYQILTAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10
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TRSEFORMTAR 51
                                                     Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PS00028; ZINC_FINGER_C2H2; 2.
553 AA; 60801 NW; EB3019D282B19092 CRC64;
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                                                                                                                           129
129 AA;
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                                                       Conservative
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                                                                                                                           14890 MW;
                                                                                                                                                                                                                                                                                                                        Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Cricetinae;
                                                                  59.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61.4%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B., Murphy
                                                    Score 34; DB Pred. No. 18; 2; Mismatches
                                                  2;
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                                                                                                                           43C825E6651EFDF3 CRC64;
                                                               DB
18;
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Q9P5K7;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-CCT-2000 (TrEMBLrel. 15, Last annotation update)
HYPOTHETICAL 32.0 KDA PROTEIN.
F27B13.170 or AnyaC29930.
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta
Magnoliophyta; eudicotyledons; core eudicots; Rosida
                      German Neurospora genome project;
Submitted (MAY-2000) to the EMBL/
EMBL; AL356172; CAB91690.1;
                                                                                                                                         Schulte U., Aign V.,
Nyakatura G., Mewes I
Submitted (MAY-2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EU Arabidopsis sequencing pr
Submitted (MAR-2004) to the
EMBL; AL050352; CAB43688 1;
EMBL; AL161575; CAB79751.1;
                                                                                                                                                                                                                                                                                                                                                                            01-OCT-2000
01-OCT-2000
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Submitted (MAY-1999) to t
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NCBI_TaxID=3702;
  SEQUENCE
                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                       Eukaryota; Fungi; Ascomycota; NCBI_TaxID=5141;
                                                                                                                                                                                                                                                                                  Neurospora crassa.
Eukaryota; Fungi;
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SEQUENCE 277 AA; 31987 MW;
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Mewes H.W., Mayer K.E
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7; Conser
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863 AA;
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(TrEMBLrel. 15, Last sequence update)
(TrEMBLrel. 15, Last annotation update)
sLUCAN 1, 3-BETA-GLUCOSIDASE PRECURSOR P
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                                                                                                                                      , Hoheisel J., Brandt P., I
H.W., Mannhaupt G.;
) to the EMBL/GenBank/DDBJ
  98928 MW;
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Pred.
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  EC64BADA28708040 CRC64;
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No.
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HSSP; P17119; 3KAR.
MGD; MGI:109234; Kif21b.
INTERPRO; IPR001680; -
INTERPRO; IPR001752; -
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01-MAY-2000
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SEQUENCE
                                                                                                                                               MEDLINE=99242603; PubMed=10225949; Marszalek J.R., Weiner J.A., Farlo
                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-OCT-2000
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                     PFAM; PF00225; kinesin; 1. PFAM; PF00400; WD40; 7.
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J. Cell B
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                                                                                                                                                                                                 NCBI_TaxID=10090;
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                                                                                                                                    'Novel dendritic kinesin
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                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity
7; Conserv
                                                                                                             on J.A., Farlow S.J., Chun J., Goldst dendritic kinesin sorting identified by different ng of two related kinesins: KIF21A and KIF21B."; Biol. 145:469-479/1999
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(TrEMBLrel.
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                                                                                                                                                                                                             (Mouse).
etazoa; Chordata;
theria; Rodentia;
                                                                                               AAF17084.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BAA97014.1;
AA; 121430
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           KINESINHEAVY
KINESIN_MOTOR_DOMAIN1; UNKNOWN_1
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59
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 34; DB
Pred. No. 1.8e
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 34;
Pred. No.
                                                                                                                                                                                                             Craniata; Vertebrata;
Sciurognathi; Muridae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                                                                                                                                                                                                                                                            PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Katoh T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C5C1B2D9173215C6
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                                                                                                                                                                                                               Muridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 1080;
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                                                                                                                                                 Goldstein
                                                                                                                                                                                                               Euteleostomi;
; Murinae; Mus
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Best Local Similarity 54.5
Matches 6; Conservative
                                                                                 Query Match 57.9%;
Best Local Similarity 70.0%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE; PS00678; WD_REPEATS; UNKNOWN_1.
PROSITE; PS50067; KINESIN_MOTOR_DOMAIN2; 1.
SEQUENCE 1668 AA; 186162 MW; 37145B2D867C9A57 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9NPW7 PRELIMINARY; PRT; 271 AA.
Q9NPW7;
Q9NPW7;
Q1-QCT-2000 (TrEMBLrel. 15, Created)
01-QCT-2000 (TrEMBLrel. 15, Last sequence update)
01-QCT-2000 (TrEMBLrel. 15, Last annotation update)
HYPOTHETICAL 30.8 KDA PROTEIN (FRAGMENT).
DKFZP762M186.
                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.

TISSUB-MELANOMA (MEWO CELL LINE);
Bloecker H., Boecher M., Brandt P., Mewes H.W., Weil B., Wiemann S.;
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ databases.

EMBL; AL359590; CAB94876.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                              Hypothetical protein:
NON_TER 1
SEQUENCE 271 AA; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 TSSEYQILTAR 11
| :||: |||:
259 TGTEYETLTAK 269
                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
1 TSSEYQILTA 10
| :|||| |
117 TRAEYQILLA 126
                                                                                                                                                                                              271 AA; 30764 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           59.6%;
54.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 34; DB 11; Length 1668; pred. No. 2.8e+02; 3; Mismatches 2; Indels
                                                                                          Score 33; DB 4; Length 271; Pred. No. 66; 1; Mismatches 2; Indels
                                                                                                                                                                                                     011097C174878478 CRC64;
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Search completed: June 4, 2001, 12:23:27 Job time: 590 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum DB :
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Match
                                                                                                                                                            100.0
89.1
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Gapop 10.0 ,
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64
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                                                    Y96183
R20811
R22550
R91438
                                                                                                                                                            W60558
Y33183
                                                                                                                          R91439
                                                                                                                                             R22549
                                                                                                                                                                                                                                                                    SUMMARIES
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                                                                                                                                                                                                              Description
             Human FCRI (CDNA C
Human FCRI (CDNA C
Human macrophage-s
                                                                 Human macrophage-s
Human macrophage-s
Human macrophage-s
                                                                                                                         Human macrophage-s
Human FcRI (cDNA c
                                                                                                                                                                           Oligopeptide from
high affinit
                                                                                                                                                              sfc-gammaRI
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	43	43	43	43	43	43	43	43	43	43	44	44	44	44	46	46	46	46	46	46	47	47	47	47	48	48	4.8	4.8	48	48	48	48	48	57
	67.2	67.2	67.2	7	67.2	7	7.	67.2	7.	67.2		68.8	8	œ.		71.9	71.9	71.9	71.9	71.9	73.4	73.4	73.4	73.4	75.0	•	٠	•	75.0	75.0	75.0	75.0	75.0	89.1
	828	789	789	764	738	738	517	398	333	272	4412	761	582	561	1018	1018	410	234	234	150	1728	1447	1447	4	1251	1242	w	1125	1018	377	374	374	374	399
	12	20	15	19	21		21	21	21	17	21	17	17	17	17.	15	12	21	19	20	12	22					19	19	18	20	19	19	19	21
	R13905	W68007	R62488	67	Y59302	Y29639	Y32390	Y96307	в12313	R92254	Y53666	R92255	R92256	R91065	R87028	R63759	R12428	Y87334	W74886	Y02365	R13144	в50693	Y33498	R68553	W37778	W52287	W37779	W52288	W06485	9	W97834	W97833	4	B43683
•																																		
																		J										٠.						
	Rabbit ATHERO-ELAM	Human soluble vasc	_	KDR protein sequen	Human VEGF recepto		Herpesvirus entry	Human IGFAM-19 imm	Human secreted pro	Neural cell adhes	Sequence gi/101742		Neural cell adhes	Rat interleukin-1	Human contactin.	1	d Fc(ga	Human signal pepti	Human secreted pro	Polypeptide ident	Deleted in Colorec	•	S S	Ω.				tus norveg	Rat contactin liga	FC	Human Fc recepto:		Human Fc receptor	Human cancer assoc

## ALIGNMENTS

RESULT W60558 08-OCT-1996; 07-OCT-1997; Synthetic. cell transformation; mutation; anti; oncogene. Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; dete Oligopeptide from extracellular domain of CD64. 18-AUG-1998 w60558; W60558 standard; peptide; 12 (UYUT-) RIJKSUNIV UTRECHT 16-APR-1998. WO9815833-A1. (first entry) 96EP-0202791. 97WO-NL00557 Ã. analysis; detection;

De Kruif CA, Logtenberg T;

WPI; 1998-240964/21.

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies identifying

Example 1; Page 29;

40pp; English

Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to

forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical of the same production of antibodies.

selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or immunofluorescent annalysis, and also to detect cell transformation caused by mutation in the contraction of the

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PF PN XXX
                                                                                                                                                                                                                                                                                                                                      RESULT
Y33183
  Baell JB, E
Maxwell KF,
                                                        11-SEP-1998;
06-FEB-1998;
                                                                                          04-FEB-1999;
                                                                                                                                                          Homo sapiens
                                                                                                                                                                                anti-inflammatory;
                                                                                                                                                                                                     drug design; therapy; inhibitor;
tissue damage; hypersensitivity;
                                   (ILEX-) ILEXUS PTY LTD
                                                                                                                12-AUG-1999
                                                                                                                                     W09940117-A1
                                                                                                                                                                                                                            atomic
                                                                                                                                                                                                                                     Fc receptor;
                                                                                                                                                                                                                                                         Human sFc-gammaRI protein fragment
                                                                                                                                                                                                                                                                                 15-NOV-1999
                                                                                                                                                                                                                                                                                                      Y33183;
                                                                                                                                                                                                                                                                                                                           Y33183 standard; Protein;
                                                                                                                                                                                            nflammatory
                                                                                                                                                                                                                                                                                                                                                 N
                                                                                                                                                                                                                            coordinate;
 Epa V, Gar
, McCarthy
                                                                                                                                                                                                                                     Fc-gammaRIIa; human; FCR;
                                                                                                                                                                                          modulator; Fc-gammaRIa;
                                                                                                                                                                                                                                                                               (first entry)
                                                       98US-0099994.
98US-0073972.
                                                                                         99WO-IB00367
           Garrett
                                                                                                                                                                                                                       nmaRIIa; human; FCR; model; three-dimension; 3 bioactive compound design; computer-assisted;
                                                                                                                                                                                immunoprotective;
  TD,
 McKenzie
                                                                                                                                                                                                                                                                                                                          261 AA:
           TPJ,
Hogarth PM,
zie IFC, Pie
                                                                                                                                                                                   rc-gammaR; Fc-epsilonR; IgG; IgE; inflammatory cell recruitment; RIa; immune function regulation;
Pietersz GA;
           Matthews
           BR;
                                                                                                                                                                                                                                  3-D;
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Three-dimensional structures and models of Fc receptors, useful in computer-assisted drug design

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see 021178) at position 25, p135 encodes a Ser residu while clone p98/X2 predicts a Thr residue. At position 58, p90

Ser residue 58, p90

Example 10;

Page 94a; 160pp; English.

New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms, etc.

WPI; 1999-539978/45

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XX PD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity; recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma RIA and models of other FcR provides a means for designing and producing compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to
                                                                                                                                                                          WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                               Rapid immunoselection cloning technique; cell surface antigen immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human macrophage-specific FcRI receptor encoded by clone p98.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                         N-PSDB; Q21179.
                                                                                                                                                                                                                                         (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                         13-JUL-1990;
                                                                                                                                                                                                                                                                                                         15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                         23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                        W09201049-A
                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R22549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially conforms to the specified atomic coordinates. Computer model images of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the FCR can be used to design bioactive chemical compounds, e.g. oligonucleotides, peptides, peptidomimetics and small organic molecules, by computer-assisted methods of drug design. Therapeutic compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 316-317; 326pp; English
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                                                                                                                                                                                                          Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        261
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                                                                                                                                                                                                                                                                         90US-0553759
                                                                                                                                                                                                                                                                                                         90WO-US04986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             245
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                                                                                                                                                                                                           Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         89.1%;
91.7%;
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Pred. No. 0.098;
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edsglyqceaat 12 EDSGLYQCEAAT 12 Query Match
Best Local Similarity 100
Matches 12; Conservative

100.0%;

Score 64; Pred. No. Mismatches

0.00033; DB 19;

Length 12; Indels

0;

0;

Gaps

0;

0;

Sequence

12 AA;

by mutation in (anti)oncogenes.

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RESULT
R91439
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Best Local S
Matches 11
                  The amino acid sequence (R91439) of human FCRI was detd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence
                                                                                                                                                                                              Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain
                                                                                                                                                                                                                                                        N-PSDB;
                                                                                                                                                                                                                                                                                                                Aruffo
                                                                                                                                                                                                                                                                                                                                                                                            13-JUL-1990;
18-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence, a single
                                                                                                                                                      Example 10; Column 55-56; 79pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapy; diagnosis;
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253 edsglywceaat 264
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DB; T14718.
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11; Conservative
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                                                                                                                                                                                                                                                                                                                Seed
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                                                                                                                                                                                                                                                                                                                                                                                        92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
93US-0139273.
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ector; FcRI; Fc receptor.
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translated
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Pred. No. 0
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1 product"
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those of the pl35
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                                                                                                                                                                                                                useful for isolation
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RESULT
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                                                                                                                                                                                                                                                                                                                                                                    panning; immunod
immune.disorder;
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                                                                                                                                                                                                                                                                                                                                                                    ge; FCRI; cell surface antigen; human; immunoselect
immunodiagnosis; diagnosis; immunotherapy; gene th
isorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 344 AA;
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9 117
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56
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140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A fusion protein in increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 344;
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                                                                                                                                                                                                                                                                                                                                                                           therapy;
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RESULT R20811 ID R3 XX AC R3

R20811 standard;

Protein; 374 AA.

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Вþ

253 edsglywceaat 264

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immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FcRI and a receptor ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
                   surfaces other than macrophages, protein of FCRI and a receptor 1 the potencies of antibodies in t
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                                                                                                                                                                                                                                                                                                                                             2000-586382/55
DB; A50631.
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88US-0160416.

89US-0379076.

90US-0498809.

90US-0553759.
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220
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268
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                  es, is useful c ligand will n therapy.
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                                                                                                   include asthma,
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(panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,
                                                                                                                                                                                                                                                                                                                                                      The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS colls using a novel method of the invention designed to isolate CSA nucleic acids. The method is based upon transient expression of a CSA in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
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Query Match Best Local S Matches 11

Local Similarity nes 11; Conserv

Conservative

89.1%;

Score 57; DB Pred. No. 0.13 0; Mismatches

.13;

1; Indels.

0;

Gaps

0;

21-MAY-1992

(first entry)

DB 21; Length 344;

1 EDSGLYQCEAAT 12

Sequence

.344 AA;

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RESULT
R22550
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Matches 11
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                                                                                                R22550;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Page 94a; 160pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-056864/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rapid immunoselection cloning technique; immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAY-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GEHO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                           253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                L5-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                           1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                        edsglywceaat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity 91.7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90WO-US04986
                                                                                                                                                                Protein;
                                                                                                                                                                                                                                                                                                                                        264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        89.1%;
91.7%;
                                                                                                                                                                374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 57; DB 13; Pred. No. 0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    technique; cell surface antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                it - for . . . neoplasms, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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RESULT
R91438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 89.:
Best Local Similarity 91.:
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts Val while clones p135 and p98/X2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a shorthydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be believed the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the protection of the 
                                                                                                                                                                                                                                                                                               Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FCRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R91438;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rapid immunoselection cloning technique; cell surface antigen; immunodiagnosis; high affinity receptor.
       Misc-difference
                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                            Human FcRI (cDNA clone pl35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-OCT-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This amino acid sequence was predicted from the cDNA sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Page 94a; 160pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human macrophage-specific FcRI receptor encoded by clone p90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            edsglywceaat 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            increase the potency of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9008-0553759
                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90WO-US04986
                                      Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      89.1%;
91.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 57; DB 13
Pred. No. 0.14;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in therapy.
                                                                                                                                                                                                                                                                                                                                  immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 374;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ig-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Вb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Š
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W00859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 89.1
Best Local Similarity 91.7
Matches 11; Conservative
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The amino acid sequence (R91438) of human FCRI was detd. from a CDNA clone, p135 (T14717), obtd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p98 (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will because it is a fixed to the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor ligand will be the first and a receptor li
                                                                                                                                                                                                                                                                                                                                                                                                                                               Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FCRI; FC receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W00859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W00859 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-200279/20
N-PSDB; T14717.
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                                                                                                                                                   25-FEB-1988;
                                                                                                                                                                                                                         09-APR-1996
                                                                                                                                                                                                                                                                                                   US5506126-A
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aruffo A, Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ||||||| ||||||
253 edsglywceaat 264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 EDSGLYQCEAAT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88US-0160416.
89US-0379076.
90US-0553759.
                                                                                                                                                   88US-0160416
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89.18;
91.78;
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Pred. No. 0.14;
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(GEHO ) GEN 13-JUL-1990; 18-OCT-1993;

HOSPITAL CORP

90US-0553759. 93US-0139273.

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Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                    Misc-difference
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                                                                                                                                                                                                                                                                 Macrophage; FcRI; cell surface antigen; panning; immunodiagnosis; diagnosis; immunodiagnosis; mindune disorder; infection; asthma; immunodiagnosis; multiple sclerosis; parasi
                                                                                                                                                                                                                                                                                                                             Human macrophage-specific FcRI
                                                                                                                                                                                                                                                                                                                                                                             Y96134;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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immune-complex disease;
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25-FEB-1988;
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23-MAR-1990;
13-JUL-1990;
                      Stamenkovic I,
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Misc-difference 140
                                      (GEHO ) GEN HOSPITAL CORP
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)B; A50592.
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Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti immune-complex disease, parasitic

Example 10; 75pp; English.

based upon transient expression of a CSA in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is animals; including proteins they encode, are useful immunotherapeutic applications, i CSA nucleic acids isolated by the method of the invention, and treatment of immune-mediated humans These eful for immunodiagnostic and ns, including the diagnosis an infections, diseases, and di e antigen by adhesion to such as a culture dish. include asthma diseases or multiple disorders in E.

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Matches 11
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                                                    cells expressing FC receptors. FogammaRI molecules are expressed by monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon FC receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant FC receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sclerosis. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 26-28; 60pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-367968/32
N-PSDB; A27466.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel Fc receptor lacking transmembrane domains, a signal peptide, and glycosylation, useful for diagnosing and treating immune disorders and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PLAC ) MAX PLANCK GES
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(MM), or rheumatoid arthritis. In addition, pharmaceutical tions containing recombinant proteins may be used to treat o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   374 AA;
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91.7%;
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            antithyroid; antiallergic; antibacterial; antiviral; dermatological; neuroprotective; cardiant; thrombolytic; coagulant; nootropic; vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating medica conditions and diagnosing pathological conditions. Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection.
                                                                                                                                                                             and cells the genes are expressed in Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory; antithyroid; antiallergic antibactorial.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammation;
immune disorder; haematopoietic cell disorder; autoimmune disorder;
allergic reaction; graft versus host disease; organ rejection;
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allergic reactions,
modulate haemostatic
                                                                                                                                                                                                                                                                                                       Claim 11;
                                                                                                                                                                                                                                                                                                                                     Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer - \,
                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; C77892.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAR-2000; 2000WO-US05882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200055350-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        dermatological; neuroprotective; thrombolytic; coagulant; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral;
antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human cancer associated protein sequence SEQ ID NO:1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B43683 standard; Protein; 399 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Rosen CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neurological disease; drug screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   haemostatic; thrombolytic; cardiovascular disorder; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              253 edsglywceaat~264
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12
                                                                                                                                                                                                                                                                                                                                                                                                           2000-587533/55.
                                                                                                                                                                                                                                                  to C78448 encode the human cancer associated proteins to B44239. The proteins can have activities based on t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer associated gene; cancer antigen; detection;
                                                                                                                                                                                                                                                                                                       Page 1739-1740;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic; proliferative; vulnerary; immunomodulator;
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91.78;
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Pred. No.
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Best Local 9
01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancers, cardiovascular disorders, neurological disease and bacterial viral infections. The peptides, nucleotides, artibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and B44240 represent sequences used in the exemplification of the present
                                                                               03-NOV-1998
                                                                                                                                         Misc-difference
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                                                         21-MAY-1997;
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                                                                                                                                                                                      Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention
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11; Conservative
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88US-0160416.
89US-0379076.
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                                                         97us-0861205
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159
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256
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171
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91.7%;
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Pred. No.
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                                                                                                                             GGG"
                                                                                                                                                                                                                        GGC"
                                                                                                                                                                                                                                               CCC.
                                                                                                                                                                                                                                                                                                                                         ccc"
                                                                                                                                                                                                  AAC"
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                                                                                                                                                                           ATG"
                                                                                                                                                                                                                                                                      CAA"
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0.15;
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has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                adnession to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a cukaryotic cell. It has been used to close the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of the coate of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression libraries is based on transient expression of an antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p135 (see V63456) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAY-1997;
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                                                                                                                                      treatment of immune-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p90 (see X07372) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in eukaryotic cells and selection of cells expressing the antigen by
                                                                                                                                                                       mmunotherapeutic applications
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1998-609251/51
DB; V63456.
                                                                                        including
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                                                                                                 humans
                                                                                                                                                                                including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           from mammalian
                                                                                                                                           disorders
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Query Match 75.0%; Score 48; DB 19; Length 374;
Best Local Similarity 83.3%; Pred. No. 3.8;
Matches 10; Conservative 0; Mismatches 2; Indels 0; Gaps

Qy 1 EDSGLYQCEAAT 12
Db 253 edsclywceaat 264

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RESULT
W97833
 Misc-difference
                                                                             cloning
                                                                                                   Human Fc receptor I
                                                                                                                 07-JUN-1999
                                                                                                                               W97833;
                                                                                                                                            W97833 standard; Protein; 374 AA.
             Misc-difference
                            Misc-difference
                                                               Homo sapiens
                                                                                    Fc receptor
                                          Misc-difference
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                                                                                    H,
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                                                                                    FCRI;
/note= "encoded by GAC" 45
                             23
                                                Location/Qualifiers
                    note= "encoded by CTC"
                                   'note=
                                                                                    cel1
                                   "encoded
                                                                                    surface antigen; lymphocyte; human;
                                   bу
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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                    expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
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N-PSDB; X07372.
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                                                                                                                                                                                                                                                                                                    induction therapy. were also obtained.
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                                                                                                                                                                                                                                                                                                  cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p98 (see X07373) were also obtained. A novel method for cloning cDNAs from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL
                                     immunotherapeutic applications,
including humans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seed
                    immune-mediated infections,
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89US-0379076.
90US-0553759.
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103 ·
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                    diseases,
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                    and disorders of
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Sequence

374 AA

WPI; 1998-609251/51. N-PSDB; X07373.

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RESULT 15
W97834
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Best Local Similarity
watches 10; Conserv
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                                          25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
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Aruffo A, Seed B;
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                                                                   01-DEC-1992;
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                 (GEHO ) GEN HOSPITAL CORP
                                   21-MAY-1997;
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                                          88US-0160416.
89US-0379076.
90US-0553759.
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77
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                                   970S-0861205
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159
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83.3%;
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New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences

Example 10; Column 53-54; 75pp; English.

An This is the amino acid sequence of human Fc receptor I (FcRI), as C deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid CC immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells CF from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372) conduction therapy. Clones p135 (see V63456) and p90 (see X07372) conduction therapy. Clones p135 (see V63456) and p90 (see X07372) conduction interleukin-2 in expression libraries is based on transient expression of an antigen construction of cells expression of an antigen by conduction of cells and selection of cells expressing the antigen by conduction and cloning of any protein which can be expressed and consistent cell surface membrane of a enkaryotic cell. It can be need to clone genes (see W63442-63) encoding cell surface can be expressed in a prokaryotic or eukaryotic host cells to consider the encoded protein. The invention also provides high cefficiency expression vectors (see W63441 and V6344) which allow the immunotheranentic and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and proteins are useful for immunodiagnostic and community and the diagnostic and community a animals, immunotherapeutic applications, including the diagr including humans. including the diagnosis and and disorders of

Sequence 374 AA;

Query Match 75.0%; Best Local Similarity 83.3%; Matches 10; Conservative 0 Score 48; DB 19; Length 374; Pred. No. 3.8; Mismatches 2; Indels 0; Gaps

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Search completed: June 4, 2001, 12:13:31 Job time: 204 sec

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Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                   Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Searched:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Run on:
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                                                                                                                                                                                                                                                                                     Score
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length: 0
length: 2000000000
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Match
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Gapop 10.0 , Gapext 0.5
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GenCore version 4.5 Copyright (c):1993 - 2000 Compugen Ltd.
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72
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              PCT-US94-05277-2
US-08-328-383B-13
US-08-375-2-307B-12
US-08-752-307B-12
US-08-768-408-093-6
US-08-714-901-6
US-08-714-901-6
US-08-828-383B-11
US-08-282-351-6
US-08-874-678-2
US-08-8874-678-2
US-08-8810-116-8
US-08-98-91-116-8
US-07-930-548A-8
US-07-930-548A-8
US-07-US93-00031-13
PCT-US93-00031-23
US-08-261-304-2
PCT-US93-00031-3
PCT-US93-00031-3
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Query Match Best Local S Matches 8

Similarity 80.88; Conservative

75.0%;

Score 48; DB Pred. No. 2.1; 1; Mismatches

DB 2; 2.1;

Length 605; Indels

0;

Gaps

0,

28 42 65.6 806 2 US-08-443-861-5 29 42 65.6 1807 1 US-07-813-393-4 31 42 65.6 1807 1 US-07-813-393-4 32 42 65.6 1307 1 US-07-977-451-6 32 42 65.6 1307 1 US-07-977-451-6 34 42 65.6 1307 1 US-07-977-451-6 34 42 65.6 1307 1 US-08-22-317-6 34 42 65.6 1307 1 US-08-22-317-6 35 42 65.6 1307 1 US-08-22-317-6 36 42 65.6 1307 1 US-08-22-317-6 37 42 65.6 1307 1 US-08-22-317-6 38 42 65.6 1307 2 US-08-23-318-2 42 43 41 64.1 1307 2 US-08-23-318-2 43 41 64.1 190 1 US-08-23-318-2 44 41 64.1 190 1 US-08-23-318-2 45 41 64.1 190 1 US-08-23-318-2 46 41 64.1 190 1 US-08-23-318-2 47 41 64.1 190 1 US-08-23-318-2 48 41 64.1 190 1 US-08-23-318-2 49 42 65.6 1367 5 PCT-US39-07610-6 49 42 65.6 1367 5 PCT-US39-07610-6 40 42 65.6 1367 5 PCT-US39-07610-6 41 42 65.6 1367 5 PCT-US39-07610-6 42 41 64.1 190 1 US-08-23-318-2 43 41 64.1 190 1 US-08-23-318-2 44 41 64.1 190 1 US-08-23-318-2 45 41 64.1 190 1 US-08-23-318-2 46 41 64.1 190 1 US-08-33-32-2 47 EXEMPTION BECAUTELY Sean A. APPLICAMT: MCCARTHY SEAT BECAUTHY SEAT A. APPLICAMT: MCCARTHY SEAT A. APPLICAMT: MCCARTHY SEAT BECAUTHY SEAT A. APPLICAMT: WESTERS FORM: APPLICAMT: WESTERS FORM: APPLICAMT: WESTERS FORM: APPLICAMTION SEATER WINDOWS SEATERS OF SEQUENCE SEATERS OF S

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303 EDEGLYECEA 312

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Query Match
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SEQUENCE CHARACTERISTICS:
LENGTH: 1018 am 'n
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
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                                                                                                                     APPLICANT:
                                                                                                                                        APPLICANT:
                                                                                                                                                       APPLICANT: Jarosz, David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                 CORRESPONDENCE ADDRESS:
                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
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                                                 NUMBER OF SEQUENCES:
                                                                     TITLE OF INVENTION
                                                                                     APPLICANT:
                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                            302 EDEGLYECEA 311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Coruzzi, Laura A. REGISTRATION NUMBER: 30,742 REFERENCE/DOCKET NUMBER: 76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 26 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/452,052
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
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E: Banner, Birch, McKie & Beckett
1001 G Street, N.W.
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                                                                 Vogelstein, Bert
Zabrecky, James R.
VENTION: Antibodies Specific for DCC Gene Product
                                                                                                                                                                   Bruskin, Arthur
                                                                                                                     Johnson, Karen
Kinzler, Kenneth W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unknown
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N: 435
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 13, Application US/08928383B Patent No. 6210921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA
                                                                                   PRIOR APPLICATION DATÁ:
APPLICATION NUMBER: US 60/026,100
FILING DATE: 13-SEP-1996
                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Robert W. Finberg, Jeffrey M. Bergelson, APPLICANT: and Marshall S. Horwitz.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                            FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                        ITLE OF INVENTION: CAR, A No. 6210921el Coxsackievirus and Adenovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    111 DEGLYOCEAS 120
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATIONS TO THE PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
                                                                                                                                      APPLICATION NUMBER: FILING DATE: 12-SEE CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                  CITY: Boston
STATE: Massa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                 NAME: Mandragouras, Amy E. REGISTRATION NUMBER: 36,2
                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                    STREET:
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                    DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible /
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                       12-SEP-1997
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Pred. No. 7
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TELEFAX: (617)742-421 NFORMATION FOR SEQ ID NO:

(617)742-4214

617)227-7400

TELEPHONE:

SEQUENCE CHARACTERISTICS

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Query Match
Best Local Similarity
Best Similarity
Best Local Similarity
Best Local Similarity
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Best Local Similarity 70.0%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             quence 94,
                                                                                                                        NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 91 amino acids
TYPE: amino acid
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FRAGMENT TYPE:
                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                      REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0 FILING DATE: 31-OCT-1994
                                                                                                              STRANDEDNESS:
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                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 54:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   48
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                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                              TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE:
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MCKENZIE, lan .
MCKENZIE, lan .
                                                                                                                                                                                              (202)672-5399
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HULETT, Mark D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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                                                                                                                                                                                                              (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                         Floppy disk
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internal
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                          71.98;
                                                                                                                                                                                                                                                                                                                                                                  US/08/332,562A
                                                                                                                                                                    94:
             Score 46; DB 2;
Pred. No. 0.63;
1; Mismatches
                                                                                                                                                                                                                                       54270/119/GRHA
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                                       DB 2; Length 91;
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             Indels
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             Gaps
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Query Match
Best Local Similarity
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                                                                                     Sequence 6,
                                                                                                     08-408-093-6
                                                     atent No. 5688916
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                             TELEX: 200154
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION: NAME: Meiklejohn, Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
              TITLE OF INVENTION:
                                            APPLICANT:
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                                                                                                                                                                388 EDSGMYQCVA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER:
                                                                                                                                                                                            1 EDSGLYQCEA 10
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                                                                                                                                                                                                                                                                                                                                                amino acid
                                                                                     Application US/08408093
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                                                                                                                                                                                                                                                                                                                                                                 607 amino acids
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                         Hemperly,
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                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IBM Compatible
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                                            Robert A:
                                                                                                                                                                                                                                         71.9%;
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Acid Sequences
            Human Cell Adhesion Molecule and Nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENCODING NOVEL SECRETED OR MEMBRANE-ASSOCIATED PROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anita L.
                                                                                                                                                                                                                                         Score 46;
Pred. No. 4
                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                       DB 2;
                                                                                                                                                                                                                                                      Length 607;
                                                                                                                                                                                                                           0; Gaps
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NAME: Fugit, Donna R.
REGIERATION NUMBER: 32,135
REFERENCE/DOCKET NUMBER: P:
NFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                  NAME/KEY: Domain LOCATION: 707..760 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                         NAME/KEY: Domain
LOCATION: 604..657
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                        NAME/KEY: Domain LOCATION: 809..857 OTHER INFORMATION: OTHER INFORMATION:
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NRIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
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                                  OTHER
                                                                                             OTHER INFORMATION:
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                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: UZIP: 07417
                                                OCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION:
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                               KEY: Domain
TION: 905..952
NIFORMATION:
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SYSTEM: PC-DOS/MS-DOS
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416..464
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138..191
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/label= FLR
/note= "conserved core of fibronectin type
III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat" <</pre>
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/note= "conserved
III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat"
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RESULT 8 US-08-408-420A-6

Sequence 6, Application US/08408420A Patent No. 5731154 GENERAL INFORMATION:

APPLICANT:

Reid, Robert A. Hemperly, John J.

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                                                          Query Match
Best Local Similarity
Matches 7; Conserv
303 EDEGIYECEA 312
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                                                                       71.98;
70.08;
                                                                                                                                            //label= ASN-glycos
/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
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glycosylation"
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/note= "potential site of ASN-linked
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/note= "potential site
                                                                      Score 46; DB.
Pred. No. 7.6;
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FEATURE:
NAME/KEY:
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NAME: Fugit, Donna R.
REGISTRATION NUMBER: 32,135
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NAME/KEY:
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OTHER INFORMATION:
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LOCATION:
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LOCATION:
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CLASSIFICATION:
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809..857
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707..760
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416..464
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332..371
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                                                                   /label= FLR
/note= "conserved
III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat"
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/note= "conserved core of fibronectin type
III-like repeat"
/label= FLR
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Best Local Similarity
Matches 7; Conserv
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70.0%;
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/note= "potential site
glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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glycosylation"
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glycosylation"
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/note= "potential site of ASN-linked
glycosylation"
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/note= "potential site of ASN-linked
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/note= "potential site of ASN-linked
                                                     Score 46; DB Pred. No. 7.6; 2; Mismatches
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7.6;
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US-08-714-901-6

Sequence 6, Application US/08714901
Patent No. 5739289
GENERAL INFORMATION:

APPLICANT:

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REFERENCE/DOCKET NUMBER: NFORMATION FOR SEQ ID NO: 6:
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MEDIUM TYPE: Floppy disk
                                                                          NAME/KEY: Domain LOCATION: 707..760 OTHER INFORMATION: OTHER INFORMATION: OTHER INFORMATION:
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APPLICATION NUMBER: US/08/040,741
FILING DATE: 26 MAR 1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                       LOCATION: 604..657
OTHER INFORMATION:
OTHER INFORMATION:
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LOCATION: 809..857
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/note= "conserved core of fibronectin type
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LOCATION: 707..760
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MEDIUM TYPE: Floppy disk
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NAME/KEY: Domain LOCATION: 809..857 OTHER INFORMATION:
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ADDRESSEE: and Company
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/note= "potential site of ASN-linked
glycosylation"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= FLR
/note= "conserved
III-like repeat"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      glycosylation"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "conserved core of fibronectin type
III-like repeat"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= ASN-glycos
/note= "potential site of ASN-linked
                              2;
                                          Score 46; DB 3; Pred. No. 7.6;
                            .Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 core of fibronectin type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of ASN-linked
                                                        Length 1018;
                              Indels
                              0;
                              Gaps
                              0,:
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303 EDEGIYECEA 312

RESULT 11 US-08-928-383B-11

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RESULT 12
US-08-282-951-6
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Patent No. 6210921
                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity Matches 6; Conser
                                                                                                                                                                                                                              Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (617)742-421
NFORMATION FOR SEQ ID NO:
                                                                                                                                        TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 72 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Morrison & Foerster
                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Mandragouras, Amy E.
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: DF
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Robert W. Finberg, Jeffrey M. Bergelson,
APPLICANT: and Marshall S. Horwitz
TITLE OF INVENTION: CAR, A No. 6210921el Coxsackievirus and Adenovirus
TITLE OF INVENTION: Receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 13-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 1
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
STATE: Massa
                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                       46 QDAGIYECESKT 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/026,100 FILING DATE: 13-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET:
                                  STATE:
                                                                                  ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                        1 EDSGLYQCEAAT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02109
94304-1018
                               Palo Alto
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28 State Street
                                                                  755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : (617)227-7400
(617)742-4214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
                                                                                                                                                                                           YANG, ZHI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LAHIVE & COCKFIELD, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide
internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                          70.3%;
                                                                                                                                        PROTEINS
                                                                                                                                                            CLONING GENES: WHICH ENCODE CELL-SURFACE AND SECRETED
                                                                                                                                                                           METHOD FOR ISOLATING AND DIRECTLY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US/08/928,383B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11:
                                                                                                                                                                                                                                                                                                                                                                                                           5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            Score 45; I
Pred: No. 0
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US-08-874-678-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                              APPLICATION NUMBER: US 08/643,839 FILING DATE: 07-MAY-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 33,943
REFERENCE/DOCKET NUMBER: 20
FELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
            NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24.190
REFERENCE/DOCKET NUMBER: A-6
FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        86 EDAGIYRCQA 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: FILING DATE: 29-JUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE:
                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: HEREW
                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1:30
                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: Four, Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 EDSGLYQCEA 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION:
                                                                                                                                                                                                                                                                                                                     94111-4187
                                                                                                                                                                                                                                                                                                                                                                      San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       70614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08874678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LEHNHARDT,
                                                                                                                                                                                                                                                                                                                                                      California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     272 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (415) 494-0792
                                                                                                                                                                                                                                                                                                                                     United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ferrara, Napoleone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chen, Helen H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Davis-Smyth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PatentIn Release #1.0, Version #1.25
(415)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Floppy disk
                                                                                                                                                                          HEREWITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUSAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                         NOVEL INHIBITORS OF VASCULAR ENDOTHELIAL GROWTH FACTOR ACTIVITY, THEIR USES AND PROCESSES FOR THEIR PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            67.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US/08/282,951
                                                                                                                                                                                            US/08/874,678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Terri L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 43;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20296-20012.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 272;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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; STRANDEDNESS:
; TOPOLOGY: unk
; MOLECULE TYPE:
US-08-643-839-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
 Best Local Similarity Matches 7; Conser
                                Query Match
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                                                                                                                                                                TELEX: 910 277299
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 767 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENERAL INFORMATION:
                                                                                                                                                                                                                             NAME: Dreger, Walter H.

REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-63291/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION: NAME: Dreger, Walter H.
                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         730 EDEGLYTCOACS 741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0 FILING DATE: 07-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Four Embarca
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RY: United States 94111-4187
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                                                                                                                                                  amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08643839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               910 277299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Flehr, Hohbach, Test, Albritton & Herbert Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               767
 67.2%;
nilarity 58.3%;
Conservative
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                                                                                                                     unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acids
                                                                                                    protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            unknown
                                                                                                                                   unknown
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58.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRODUCTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOVEL INHIBITORS OF VASCULAR ENDOTHELIAL GROWTH FACTOR ACTIVITY, THEIR USES AND PROCESSES FOR THEIR
                                                                                                                                                                                                                                                                                                                                                                                   US/08/643,839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Terri L
Score 43; DB Pred. No. 17; 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 43;
Pred No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                  DB 3;
 3; Indels
                                  Length 767;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 767;
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 0;
 Gaps
 0;
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EDSGLYQCEAAT 12

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Query Match
Best Local Similarity
"---hes 6; Conserv
                                                                                       밁
                                                                                                                    Qy
Search completed: June 4, 2001, 12:14:31 Job time: 224 sec
                                                                                                                                                                                                                                            ; MOLECULE TYPE: US-08-261-304-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-261-304-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/08261304
                                                                                                                                                                                                                                                                                                                                TELEPHONE: (202) 833-7
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                   NAME: POTTER, Jane E. R. REGISTRATION NUMBER: 33,332 REFERENCE/DOCKET NUMBER: 06' TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 833-7533
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/649,565
FILING DATE: 01-FEB-1991
APPLICATION NUMBER: U.S. 07/487,038
FILING DATE: 02-MAR-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: ASCII CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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                                                                                       661 EDAGVYECES 670
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 1225 Conne
STREET: Suite 300
                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                      1 EDSGLYQCEA 10
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: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Sterne, Kessler, Goldstein & Fox 1225 Connecticut Avenue, N.W.
                                                                                                                                                                                                                                                                                                                    828 amino acids
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                                                                                                                                                          Conservative
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                                                                                                                                                                         Score 43; DB 1;
Pred. No. 19;
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GenCore version 4.5 Copyright (c) 1993 - 2000 Comp

Compugen Ltd

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 s
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12.645 Million cell updates/sec
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R;Allen, J.M.; Seed, B.
Science 243, 378-381, 1989
Science 243, 378-381, 1989
A;Title: Isolation and expression of functional high-affinity Fc receptor complementa A;Reference number: A41357; MUID:89100284

A; Accession: A41357

C;Date: 03-Apr-1992 #sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000 C;Accession: A41357; S03019

N; Alternate

Species: Homo sapiens (man)

names: CD64

gamma (IgG) receptor I (high affinity) form b -

human

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ALIGNMENTS										•						
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Fc gamma (IGG) receptor I-A (high affinity) precursor - human
N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000
C;Accession: A39878; I7/304; B41357; S03018; I57525
C;Accession: A39878; I7/304; B41357; S03018; I57525
R;van.de Winkel, JG.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M.
J; Biol. Chem. 266, 13449-13455, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 A;Note: the authors translated the codon ACT for residue 25cas Ala C;Superfamily: Fc gamma receptor 1; immunoglobulin homology C;Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein F;117-170/Domain: immunoglobulin homology <IMM>
A;Title: Gene organization of the human high affinity receptor for IgG, A;Reference number: A39878; MUID:91302383
A;Accession: A39878
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A39878
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334 R;Allen, J.M.; Seed, B. McLeic Acids Res. 16, 11824, 1988
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A; Residues: 1-344 <ALL2>
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A; Residues: 1-344 <ALL1>
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Best Local :
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91.7%;
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A; Molecule type: DNA A; Residues: 1-374 < VAN>

nces: EMBL:U55211; NID:g1305526; PID:g1305527; PIDN:AAA99159.1 neural cell adhesion molecule L1; fibronectin type III repeat

III repeat homology

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A; Molecule type: mRNA
A; Residues: 1-24,'5', 26-337,'T', 339-374 <AL2>
A; Residues: 1-24,'5', 26-337,'T', 339-374 <AL2>
A; Residues: 1-24,'5', 26-337,'T', 339-374 <AL2>
A; Cross-references: EMBL:X14356; NID::g31331; PIDN:CAA32537.1;
A; Cross-references: Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compared to Microsoft State of the Compar
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A; Accession: B41357
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                                                                                                                                                           submitted to the EMBL Data A; Description: Molecular cl
                                                                                                                                                                                                                                                                                            C;Date:
                                                                                                                                                                                                                                                                                                                                                     neural cell adhesion molecule L1 - N;Alternate names: E587 antigen
                                                                A; Status:
                                                                                              A; Accession: T43027
                                                                                                                                A; Reference number:
                                                                                                                                                                                                                           R;Giordano,
                                                                                                                                                                                                                                                          C; Accession:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cross-references: GB:L03418; NID:g184840; PIDN:AAA36049.1; PID:g292169;Allen, J.M.; Seed, B.
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Accession: 157525
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Title: Nucleotide sequence of three cDNAs
Reference number: S03018; MUID:89098339
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   Residues:
                                                                                                                                                                                                                                                                                                                          Species: Carassius auratus (goldfish)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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59,78,152,159,163,195,240/Binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Superfamily: Fc gamma
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Residues: 1-8 <RE2>
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Residues: 1-24,'S',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           117-170/Domain: immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cross-references: GB:S57204; NID:g298692; PIDN:AAD13887.1; PID:g4261587
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coss-references: GB:X14356; GB:M21091; NID:g31331; PIDN:CAA32537.1; PID:g31332
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                                                            preliminary; translated
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                                                                                                                                                                                                                           S.; Laessing,
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î-1232 <GIO>
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                                                                                                                                                                                                                                                                                        #sequence_revision 11-Jan-2000 #text_change
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                                                                                                                                                           , U.; Lottspeich, F.; Stuermer, C.A. ta Library, April 1996 cloning of goldfish E587 antigen, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89.1%;
91.7%;
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ding site: carbohydrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 57; DB Pred. No. 0.01 0; Mismatches
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                                                                   from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                          goldfish
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lycoprotein; immunoglobulin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene products
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0.018;
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                                                                                                                                                                     adhesion molecule expr
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C;Superfamily: neura
C;Keywords: cell adl
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S05944
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A;Cross-references: GB:M81846; NID:g212206; PIDN:AAA48924.1; A;Experimental source: embryo C;Superfamily: interleukin-1 receptor type I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gene 111, 239-243, 1992
A;Title: The chicken II-1 receptor: differential evolution
A:Reference number: JQ1526; MUID:92175529
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            interleukin-1 receptor I precursor - chicken
C; Species: Gallus gallus (chicken)
C; Date: 17-Aug-1992 #sequence_revision 17-Aug-1992 #text_change 29-Sep-1999
                                                                                                                                                                                                                                                                                                                                neuronal cell surface protein F3 precursor - mouse C;Species: Mus musculus (house mouse) C;Deceles: Mus musculus (house mouse) C;Date: 01-Dec-1989 #sequence_revision 01-Dec-1989 #text_change C;Accession: S05944 R;Gennarini, G; Cibelli, G; Rougon, G; Mattei, M.G.; Goridis, R;Gennarini, G; Cibelli, G; Rougon, G; Mattei, M.G.; Goridis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Keywords: cytokine receptor; transmembrane protein F;I-19/Domain: signal sequence #status-predicted <SIG> F;20-555/Product: interleukin-1 receptor I #status predicted <INT> F;339-359/Domain: transmembrane #status predicted <TRA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R; Guida, S.
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                                                                                               A; Map position: 15F
C;Superfamily: contactin; fibronectin type III repeat homology;
F;1-20/Domain: signal sequence #status predicted <SIG>
F;256-312/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-1020 <GEN>
A; Cross-references: EMBL:X14943;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F;339-359/Domain: transmembrane #status predicted
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                                                                                                                                                                           C; Genetics:
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                                                                                                                                                                                                                                                        A; Accession: S05944
                                                                                                                                                                                                                                                                        A; Reference number: S05944;
                                                                                                                                                                                                                                                                                               A;Title: The mouse neuronal
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                Query Match
Best Local
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nes 8; Conser
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8; Conser
Similarity
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88.9%;
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66.7%;
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Score 48;
Pred. No.
1; Mismatc
                                                                                                                                                                                                NID: g50937; PIDN: CAA33075.1;
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Pred. No. 0.93
1; Mismatches
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Pred. No. 1.3;
2; Mismatches
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0.93;
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                                         Length 1020
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EDEGLYECEA EDSGLYQCEA 10

312

Conservative

Mismatches

Indels

Gaps

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A; Title: The carbonic anhydrase domain of A; Reference number: A57112; MUID:95354206 A; Accession: A57112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Status: preliminary; nucleic acid sequence not shown; not compared with conceptual tr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Peles, E.; Nativ, M.; Campbell, P.L.; Sakurai,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              contactin precursor - rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: EMBL:Y00813, NID:g63328; PIDN:CAA68753.1; Note: part of this sequence, including the amino end of the Bruemmendorf, T.; Wolff, J.M.; Frank, R.; Rathjen, F.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;Ranscht,
                                                                                                                                                                                                                                                    Superfamily: contactin; fibronectin type III repeat homology.immunoglobulin homology Keywords: blocked carboxyl end; glycoprotein; lipoprotein; phosphatidylinositol linka: 1-20/Domain: signal sequence #status predicted <SIG>
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Title: Sequence of contactin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Alternate
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                                                                                                                                                                      983-1002/Domain: transmembrane #status predicted <TMM>
                                                                                                                                                                                                                                                                                                               Comment: F11 is a chick neural cell surface-associated glycoprotein implicated in neu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Species: Gallus gallus (chicken)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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Residues: 1-1021 <F
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    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Accession: JU0094
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   uron 2, 1351–1361, 1989
Title: Neural cell recognition molecule F11; homology with fibronectin type III and Reference number: JU0094; MUID:90180453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecule type: mRNA
Residues: 1-1091 <RAN>
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                                                                                                                                                  003-1091/Domain:
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ross-references: GB:X14877; NID:g1708784; PIDN:CAA33018.1; PID:g63385
ote: the carboxy-end hydrophobic stretch is compatible with the consensus motif for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tactin
                                                                                                                                                                                                             982/Domain: extracellular #status predicted <EXT>
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                                                                                                                          1091/Domain: intracellular #status predicted <INT> 249,329,448,464,485,512,582,621,924,Binding site: c
                                                                                                                                                                                                                                   091/Product: contactin #status predicted <MAT>
                        Local
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                                                                                                                                                                                                                                                                                                                                                         fll comprises six domains related to the immunoglobulin domain type C and four
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                                                                                                                                                                                           Domain: immunoglobulin
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    8
                          Similarity
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8; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1561-1573, 1988
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                   75.0%;
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80.0%;
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                                                                                                                                                                                           homology <IMM>
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Pred. No.
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Pred.
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      Mismatches
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                                         Length 1091;
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                                                                                                                                                                                                                                                                                                                                                                                                       Best
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N; Alternate names: pE4 protein
C; Species: Rattus norvegicus (Norway rat)
C; Date: 02-Aug-1994 #sequence_revision 02
C; Accession: A54017; A61206
F; Chadeneau, C.; LeMoollac, B.; Denis, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   probable tumor suppressor - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 18-Oct-1991 #sequence_revision 18-Oct-1991 #text_change 16-Feb-1997
C;Accession: A38442
R;Nigro, J.M.; Cho, K.R.; Fearon, E.R.; Kern, S.E.; Ruppert, J.M.; Oliner, Cell 64, 607-613, 1991
                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-267 <NIG>
A;Cross-references: GB:M63
C;Keywords: transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                            A; Status: preliminary A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: A38442; A; Accession: A38442
A; Title: A novel member of the immunoglobulin gene superfamily expressed in rat carci
                                                                                                                         colon carcinoma-associated antigen pE4 precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Title: Scrambled exons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Residues: 1-1256 < KRA>
A; Cross-references: EMBL: AF004840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the EMBL Data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R; Krauss,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Accession:
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                    J. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Experimental source:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Status: preliminary; translated
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Date: 24-Mar-1999 #sequence_revision
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Best Local
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8; Conserv
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9; Conser
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                    269,
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                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                         GB:M63696; GB:M63700;
                  15601-15605,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MUID: 91121517
                                   Denis, M.G
                    1994
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                                                                                                                                                                                                                                                                                                    Score 47;
Pred. No.
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September 1998
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                                                                      02-Aug-1994 #text_change 01-Dec-1995
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0.7
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A; Molecule type: mRNA
A; Residues: 1-1447 <HED>
                                                                                A;Title: The DCC gene product in cellular differentiation A;Reference number: A54100; MUID:95011532 A;Accession: A54100
                                                                                                                                          R; Hedrick, L.; Cho, K.R.; Fear
Genes Dev. 8, 1174-1183, 1994
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A;Residues: 34-41,'X',43-53 <CH2>
A;Residues: 34-41,'X',43-53 <CH2>
A;Residues: 34-41,'X',43-53 <CH2>
C;Comment: This 66K protein is detected by monoclonal antibody E4 on colon carcinoma cel
C;Keywords: glycoprotein; membrane protein
                                                                                                                                                                                                                                                    N; Alternate names:
                                                                                                                                                                                                                                                                          tumor suppressor protein DCC precursor -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Superfamily: neural cell adhesion molecule L1; fibronectin type
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A;Cross-references: EMBL:X76132; NID:g453209; PIDN:CAA53735.1; PID:g453210
R;Fearon, E.R.; Cho, K.R.; Nigro, J.M.; Kern, S.E.; Simons, J.W.; Ruppert, J.M.; Hamiltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Note: C18F3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Map position:
A; Introns: 34/3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;Chadeneau, C.; Denis, M.G.;
nt. J. Cancer 47, 903-908, 1
;Title: Characterization, is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;Residues:
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                                                                                                                                                                                        Accession: A54100; A40098
                                                                                                                                                                                                             Date: 02-Aug-1994 #sequence_revision 02-Aug-1994 #text_change 05-Nov-1999
                                                                                                                                                                                                                                 Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local :
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Cross-references: EMBL:U50067; PIDN:AAA93439.2
Experimental source: strain Bristol N2
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Accession: T30988
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Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text_change 17-Mar-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pothetical protein C18F3.2 - Caenorhabditis elegans
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to the EMBL Data Library, August 1999
tion: The sequence of C. elegans cosmid C18F3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      preliminary; translated from GB/EMBL/DDBJ
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8; Conser
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; 72/1; 131/2; 157/1;
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                                                                                                                                                                                                                                                    colorectal cancer suppressor
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                                                                                                                                                                  Fearon, E.R.; Wu, T.C.; Kinzler, K.W..; Vogelstein,
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Pred. No. 1.1;
3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 47;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             254/3; 303/1; 400/1; 495/1; 592/1;
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1.1;
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                                                                                                                             colorectal tumorigenesis.
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                                                                                                                                                                    A; Title: Structure and mapping of the gene A; Reference number: A46480; MUID:92166399
                                                                                                                                                                                                                                                                          C;Species: Mus musculus (house mouse)
C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                 망
    A; Title:
                       J. Immunol
                                                                                      A; Residues: 1-404 <OSM>
                                                                                                                           A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                     N;Alternate
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                                         R; Sears,
                                                              A; Note: sequence extracted
                                                                                                          A; Molecule type: DNA
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D.W.; Osman, N

371-378, 1990

and

expression of the mouse high affinity Fc receptor for

Tate,

from

NCBI backbone (NCBIN:85205, NCBIN:85208, B.; McKenzie, I.F.C.; Hogarth, P.M.

NCBIN: 85212

148, 1570-1575,

1992

C.A.;

McKenzie, I.F.; Hogarth,

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21-Jan-2000

encoding mouse

high affinity

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gamma

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names: high

affinity IgG receptor

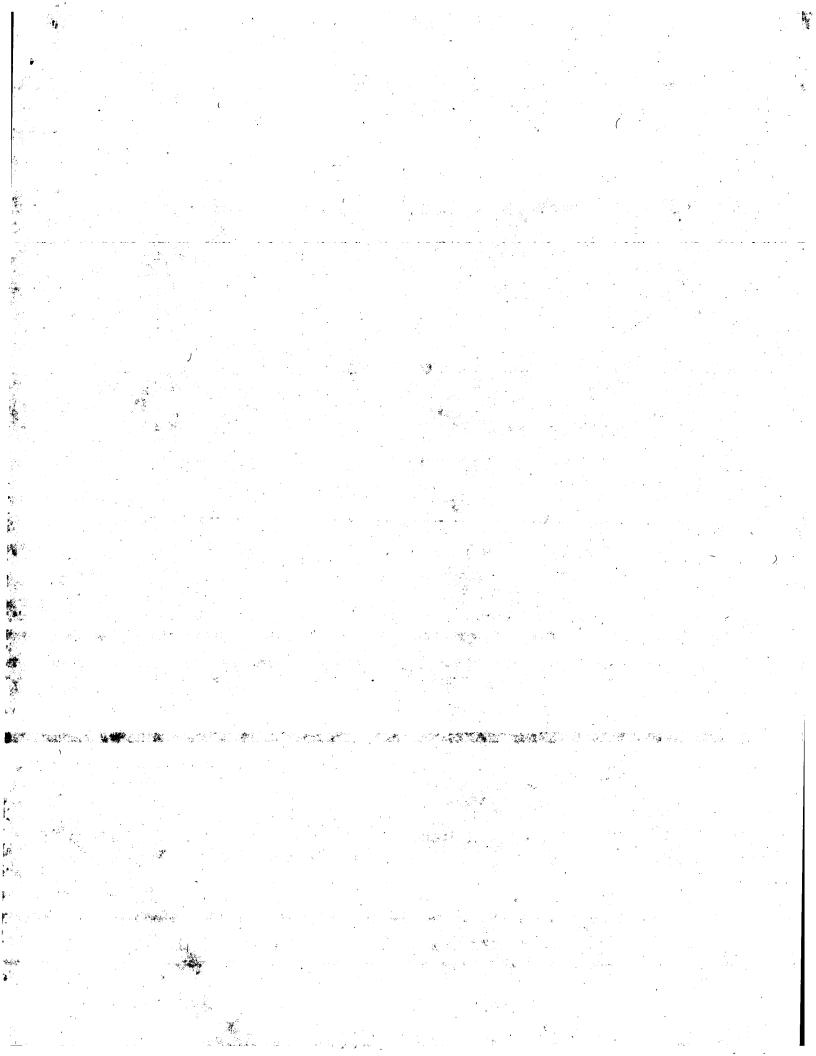
mouse

EDAGFYWCEVAT

278

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R;Prins, J.B.; Todd, J.A.; Kourryues, M.N., Chem. Science 260, 695-698, 1993
A;Title: Linkage on chromosome 3 of autoimmune diabetes and A;Reference number: I48471; MUID:93242399
                                                                                                                                                                                                                                                                                                                                                      Fc gamma (IgC) receptor high affinity - mouse
C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
                                                                                                                       A;Cross-references: EMBL:X70980; NID:g311748; PIDN:CAA50311.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: immunoglobulin receptor F;128-180/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                              A; Molecule type: mRNA
A; Residues: 1-336 < RES>
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A; Residues: 1-750 <FEA>
A; Cross-references: GB:M32292; NID:g181492;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Science 247, 49-56, 1990
A; Title: Identification of
                                                                                                                                                                                                                                             A; Accession: I48471
                                                                                                                                                                                                                                                                                                                                         C; Accession:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C;Keywords: transmembrane protein; tumor suppressor F;1-25/Domain: signal sequence #status predicted <SIG>
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A;Map position: 18q21.1-18q21.1
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Search completed: June 4, 2001, 12:15:44
Job time: 277 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neural adhesion protein F3 precursor - bovine
C;Species: Bos primigenius taurus (cattle)
C;Date: 14.Nov.1995 #sequence_revision 08-Feb-1996 #text_change 20-Jun-2000
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A;Accession: A43511
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Best Local Similarity 70.0
Matches 7; Conservative
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;Watanabe, K.;
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Best Local Similarity 66.7%;
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Status: preliminary
;Molecule type: mRNA
;Residues: 1-404 <SEA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ne 160, 245-248, 1995
Title: Cloning of the cDNA encoding neural adhesion molecule F3 from bovine brain.
Reference number: JC4211; MUID:95369697
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Residues: 1-1018 <WAT>
Cross-references: DDBJ:D32135; NID:g1060860; PIDN:BAA06861.1; PID:g1060861
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303 EDEGIYECEA 312
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245-248, 1995
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Q06805	P55068	Q61361	P21802	P18460	095025	P79280	Q28343	Q61282	P07897	Q28062	P21804
bos taurus	rattus norv	mus musculu	homo sapien	gallus gall	homo sapien	sus scrofa	canis famil	mus musculu	rattus norv	bos taurus	gallus gall

## ALIGNMENTS

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T O1-MAR-1989 (Rel. 10, Created)

T O1-APR-1990 (Rel. 14, Last sequence updated)

DT 15-UTL-1999 (Rel. 38, Last annotation upon the contactin PRECURSOR (NEURAL CELL RECOGN)

OS Gallus gallus (Chicken).

OS Gallus gallus (Chicken).

OS Gallus gallus (Chicken).

Aves; Neognathae; Gallifo
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Best Local S
Matches 11
                                                         Ranscht B., Dours M.T.;
"Sequence of contactin, a 130-kD glycoprotein concentrated in areas of interneuronal contact, defines a new member of the immunoglobulin supergene family in the nervous system.";
J. Cell Biol. 107:1561-1573(1988).
                MEDLINE=89286606;
Wolff J.M., Bruemn
                                                    GPI-ANCHOR
                                                                                                                                                                                                                           SEQUENCE FROM N.A.

MEDLINE-90180453; PubMed-2627374;

Bruemmendorf T., Wolff J.M., Rainer F., Rathjer F.G.;

"Neural cell recognition molecule F11: homology with fibronectin III and immunoglobulin type C domains.";

Neuron 2:1351-1361(1989).
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MEDLINE=89008597; Pub
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11; Conserv
          36606; PubMed=2735929;
Bruemmendorf T., Rath
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IG-LIKE C2-TYPE DOWAIN.
IG-LIKE C2-TYPE DOWAIN.
BY SIMILARITY.
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InterPro; IPR003006;
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PIR; S01998; S01998.
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-!- SIMILARITY: CONTAINS 4 FIBRONECTIN TYPE III-LIKE DOMAINS.
-!- CAUTION: REF.2 SEQUENCE DIFFERS FROM THAT SHOWN IN THE C-TE
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MEDLINE=89340657; PubMed=2474555;
Gennarini G., Cibelli G., Rougon G., Mattei M.-G., Goridis C.;
"The mouse neuronal cell surface protein F3: a phosphatidylinositol-anchored member of the immunoglobulin superfamily related to chicken contactin.";
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Mammalia; Eutheria;
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-I- FUNCTION: MEDIATES CELL SURFACE INTERACTIONS DURING NERVOUS
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P43146;
01-NOV-1995
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                                                                                                                                                  human esophageal squamous metastasis.";
                                                                                                                                                                                          Miyake S., Nagai K.,
                                                                                                                                                                                                                                                                               Cho K.R., Oliner J.D., Simons J.W., Hedrick L:, Fearon E.R., Preisinger A.C., Hedge P., Silverman G.A., Vogelstein B.;
                                                                                                                                                                                                                                                                                                                                            Nigro J.M., Cho K.R., Feard Oliner J.D., Kinzler K.W., "Scrambled exons."; Cell 64:607-613(1991).
                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE OF 107-472 FROM N.A. (SC
MEDLINE-91121517; PubMed-1991322;
                                                                                                                                                                                                                                                                                                                                                                                                                                              "Identification of a chromosome colorectal cancers.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90100559; PubMed=2294591;
Fearon E.R., Cho K.R., Nigro J.M.
Ruppert J.M., Hamilton S.R., Prei
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Mammalia; Eutheria;
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01-OCT-2000
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MEDLINE-94243823; PubMed=8
                                                                                                                                                                                                                                                                                                                  GENE STRUCTURE,
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FUNCTION: IMPLICATED AS A TUMOR SUPPRESSOR GENE.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN
TISSUE SEBCIFICITY: FOUND IN AXONS OF THE CENTRAL AND PERIPP
NERVOUS SYSTEM AND IN DIFFERENTIATED CELL TYPES OF THE INTE:
DISEASE: COLORECTAL TUMORS THAT LOST THEIR CAPACITY TO
DIFFERENTIATE INTO MUCUS PRODUCING CELLS UNIFORMLY LACK DCC
EXPRESSION, INACTIVATION OF DCC DUE TO ALLELIC DELETION AND
POINT MUTATIONS MAY CAUSE BOTH LYMPHATIC AND HEMATOGENOUS
METASTASIS OF DESOPHAGEAL SQUAMOUS CELL CARCINOMAS.
SIMILARITY: CONTAINS 6 FIBRONECTIN TYPE III-LIKE DOMAINS.
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8; Conserv
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K., Yoshino K., Oto M.,
and allelic deletion of
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Vogelstein B.;
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tumor suppressor gene DCC
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Thomas G., Kinzler K.W.
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L; M32290; AAA52176.1;
L; M63696; AAA52177.1;
L; M63700; AAA52179.1;
L; M63718; AAA52180.1;
L; M63718; AAA52180.1;
L; M63698; AAA52181.1;
               Similarity
8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               an email to license@isb-sib.ch)
                                                                                             1375
               Conservative
                                                                                                             201
                                                                                                                                                                                                                               525
619
722
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                                                                                            1-375
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                                                             421
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329
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                                                                                                                           168
                      73.4%;
                                                      158456
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            Score 47; DB 1;
Pred. No. 0.51;
1; Mismatches
                                                            MISSING
                                                                            /FTId=VAR_
                                                                                                                                                                              BY SIMILARITY.
BY SIMILARITY.
                                                                                                                           N-LINKED
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IG-LIKE C2-TYPE I
IG-LIKE C2-TYPE I
IG-LIKE C2-TYPE I
                                                                                 /FTId=VAR_003910.
P -> H (IN A COLORECTAL
/FTId=VAR_003911.
                                                                                                                                                                                                                                                                        CYTOPLASMIC (POTENTIAL).
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
                                                                                                                                                             N-LINKED (GLCNAC...
                                                                                                                                                                                                                                                                                                                                       domain;
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BY SIMILARITY.
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SING (IN REF. 3).
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                            Length 1447;
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             Indels
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(POTENTIAL)
(POTENTIAL)
                                                                                                                        CARCINOMA)
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                                                                                          CARCINOMA)
            0;
           Gaps
            0
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RESULT 5
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MGD; MGI:94869; Dcc.
InterPro; IPR001777; -
InterPro; IPR003006; -
  INIT_MET
DOMAIN
TRANSMEM
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                                                                                                                                                                                                                                                                        use by non-profit institutions as zony usage by and for commercial modified and this statement is not removed. Usage by and for commercial modified and this statement is not removed. Usage by and for commercial modified and this statement is not removed. Usage by and for commercial modified and this statement is not removed.
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01-NOV-1997
30-MAY-2000
                                                                                                                        SIGNAL
                                                                                                                                   Glycoprotein; Immunoglobulin
Anti-oncogene; Alternative in
                                                                                                                                                               PRINTS;
                                                                                                                                                                            Pfam; PF00047;
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"Cloning of the mouse homologue of gene (mDCC) and its expression in to concogene 11:2243-2254(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=BALB/C; TISSUE=Brain;
MEDLINE=96112625; PubMed=8570174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota;
Mammalia; J
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                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE SPECIFICITY: IN THE EMBRYO, EXPRESSED AT HIGH LEVELS DEVELOPING BRAIN AND NEURAL TUBE. IN ADULT, HIGHLY EXPRESSED BRAIN WITH VERY LOW LEVELS FOUND IN TESTIS, HEART AND THYMUS. DEVELOPMENTAL STACE: LOW LEVELS IN EARLY GESTATION. HIGHEST I EXPRESSED DURING MID GESTATION. LEVELS DECREASE IN LATE GESTAND REMAIN AT THIS LEVEL IN THE ADULT.
                                                                                                                                                                                                                                                                                                                                 European Bioinformatics Institute.
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(Rel. 35, Last seq
(Rel. 39, Last ann
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39, Last annotation update)
PROTEIN DCC PRECURSOR.
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Rodentia;
                                                                                                                                                                                                                                                                                                                  ormatics Institute. There are no restinstitutions as long as its content
                                                                                                                               initiation; Alternative splicing.
EXTRACELLULAR (POTENTIAL)
POTENTIAL.
CYTOPLASMIC (POTENTIAL).
IG-LIKE C2-TYPE DOMAIN.
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                                                                   ISOFORM
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                                                                                             ISOFORM
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in the developing mouse embryo.";
                                                                            SUPPRESSOR PROTEIN DCC,
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FCG1_MOUSE
P26151;
01-MAY-1992
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            between the Swiss Institute of Bioinf the European Bioinformatics Institute use by non-profit institute.
                                                   This SWI
                                                                                                                                                                                                       SEQUENCE FROM N.A.

MEDLINE=92166399; PubMed=1531670;
Osman N., Kozak C.A., McKenzie I.F., Hogarth P.M.;
"Structure and mapping of the gene encoding mouse high affinity
gamma RI and chromosomal location of the human Fc gamma RI gene.
J. Immunol. 148.1570-1575(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-MAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence up
15-JUL-1999 (Rel. 38, Last annotation
HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC
RI) (FCRI) (IGG FC RECEPTOR I).
                                                                                                                                                                                                                                                                                                                                                    MEDLINE-90111035; PubMed-2136886;
Sears D.W., Osman N., Tate B., McKenzie
"Molecular cloning and expression of the
receptor for IgG.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
Eukaryota; Metazoa; C
Mammalia; Eutheria; R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CARBOHYD
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                                                                                                 FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS ARFINITY RECEPTOR.

SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.

TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.

SIMILARITY: CONTAINS 3 IMMUNOGLOBULIN-LIKE C2-TYPE D
CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-10 IS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DEGLYQCEAS
                                                                 SWISS-PROT entry is copyright.
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7420
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Rodentia;
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80.08;
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Pred. No. 0.51;
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FIBRONECTIN TYPE-III.
FIBRONECTIN TYPE-III.
FIBRONECTIN TYPE-III.
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
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CARBOHYD
CONT_HUMAN STANDARD: PRT; 1018 AA. 012860; 012861; 014030; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 01-OCT-2000 (Rel. 40, Last annotation update) CONTACTIN PRECURSOR (GLYCOPROTEIN GP135).
                                                                                                                                                                                                                                                                                                        _HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOMAIN
                                                                                                                                   MEDLINE=95048335; PubMed=7959734;
Berglund E.O., Ranscht B.;
"Molecular cloning and in situ localization gene (CNTNL) on chromosome 12q11-q12.";
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DOMAIN
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                                                                       molecule contactin.
                                                                                Reid R.A., Hemperly J.J.; "Identification and characterization
                                                                                                 SEQUENCE FROM N.A., AND CHARACTERIZATION. MEDLINE-94217459; PubMed-8164510;
                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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PIR; A43511; A43511.
                                                                                                                              Genomics
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01-DEC-1992 (Rel. 24, Las.
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PIR; S22128; S22128.
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"The axonally secreted cell adhesion molecule, axonin-1. Primary structure, immunoglobulin-like and fibronectin-type-III-like domains and glycosyl-phosphatidylinositol anchorage.";
Eur. J. Biochem. 204:453-463(1992).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.,
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-i- FUNCTION: AXON-ASSOCIATED CELL ADHESION MOLECULE (2
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SIMILARITY: CONTAINS
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SUBCELLULAR LOCATION: ATTACHED TO THE NEURONAL GPI-ANCHOR.
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                                                 Signal; GPI-anchor;
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MEDIJNE-94140354; PubMed-8307567;
TSiotra C.P., Karagogeos D., Theodorakis K., Michaelidis M. Modi W.S., Furley J.A., Jessel M.T., Papamatheakis J.;
Modi W.S., Furley J.A., Jessel M.T., Papamatheakis J.;
"Isolation of the cDNA and chromosomal localization of the
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Homo sapiens (Homan).
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InterPro; IPR003006; -.
Pfam; PF00041; fn3; 4.
Pfam; PF00047; 19; 6.
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EMBL; X67734; CAA47963.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TAX1) encoding the human axonal glycoprotein TAG-1. Genomics 18:562-567(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-93145965; PubMed-8425542; Hasler T.H., Rader C., Stoeckli E.T., "CDNA cloning, structural features, ar human TAG-1/axonin-1.";
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SUBCELLULAR LOCATION: ATTACHED TO THE NEURONAL MEMBRANE-BY
GPI-ANCHOR AND IS ALSO RELEASED FROM NEURONS.
SIMILARITY: CONTAINS 6 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
SIMILARITY: CONTAINS 4 FIBRONECTIN TYPE III-LIKE DOMAINS.
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FIBRONECTIN TYPE-III.
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FIBRONECTIN TYPE-III.
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InterPro; IPR001
Pfam; PF00041; 1
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P22063;
01-AUG-1991 (Rel. 19, Created).
01-AUG-1991 (Rel. 19, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
15-JUL-1999 (Rel. 38, Last annotation TAG-1).
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                                                                                        use by non-profit institutions as long modified and this statement is not removed entities requires a license agreement (See
                                                                                                                       between the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There
                                                                                                                                                                                                                                                                     "The axonal glycoprotein TAG-1 is an immunoglobulin member with neurite outgrowth-promoting activity."; Cell 61:157-170(1990).
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Mammalia; Eutheria;
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                                                                                                                                                                         FUNCTION: MAY PLAY A ROLE IN THE INITIAL GROWTH AND GUIDANCE AXONS. MAY BE INVOLVED IN CELL ADHESION.

SUBCELLULAR LOCATION: ATTACHED TO THE NEURONAL MEMBRANE BY A GPI-ANCHOR AND IS ALSO RELEASED FROM NEURONS.

TISSUE SPECIFICITY: IN NEURAL TISSUES IN EMBRYOS, AND IN ADUL BRAIN, SPINAL CORD AND CEREBELLUM.

DEVELOPMENTAL STAGE: TRANSIENTLY EXPRESSED ON A SUBSET OF AXO IN THE DEVELOPING RAT NERVOUS SYSTEM.

SIMILARITY: CONTAINS 6 IMMUNOCLOBULIN-LIKE C2-TYPE DOMAINS.
                                                                                                                                                                 SIMILARITY: CONTAINS 6 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS SIMILARITY: CONTAINS 4 FIBRONECTIN TYPE III-LIKE DOMAINS.
                                                                                                                                            SWISS-PROT entry is copyright. It is produced through a collaboration
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                                                                                an email to license@isb-sib.ch).
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Rodentia;
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MEDLINE-93246254; PubMed-7683304;
Araki M., Araki K., Vassalli P.;
"Cloning and sequencing of mouse '
Gene 126:261-264(1993)
 STRAIN-129; TISSUE-Embryo;
MEDLINE-94117008; PubMed-7507076;
Cybulsky M.I., Allan Motamed M., (
"Structure of the murine VCAM1 gen
                                                                                                                                                                                                                                Hession C., Moy P., Tizard R., Chisholm P. Burkly L., Miyake K., Kincade P., Lobb R., "Cloning of murine and rat vascular cell a
                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
STRAIN-FVB; TISSUE-Lung;
MEDLINE-92181437; PubMed-1371918;
Hession C., Moy P., Tizard R., Ch
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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Biochem. Biophys. Res.
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murine VCAM1 gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Korenaga R., Ando J., Tsuboi H., Kamiya A.;
Submitted (DEC-1995) to the EMBL/GenBank/DDBJ databases:
-!- FUNCTION: IMPORTANT IN CELL-CELL RECOGNITION. APPEARS TO FUNCTION
IN LEUKOCYTE-ENDOTHELIAL CELL ADHESION. INTERACTS WITH THE BETA-1
                                                                                                                                                                                                                                                                             -!- SIMILARITY: CONTAINS 7 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  adhesion molecule (VCAM) mRNA encoding a glycosylphosphatidylinositol-anchored VCAM protein."; Proc. Natl. Acad. Sci. U.S.A. 90:5919-5923(1993).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE OF 311-345 FROM N.A. (GPI-ANCHORED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 1-693 FROM N.A. STRAIN-NIH SWISS, AND 129/SV; KUMAR A.G., Dai Y.X., KOZAK C.A., Ballantyne C.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Terry R.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=95015899;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=C57BL/6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 1-345 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Moy P., Lobb R., Tizard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 1-345 FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytokine induction of an alternatively spliced murine dhesion molecule (VCAM) mRNA encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _Cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WEDLINE=93232042; PubMed=7682556;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISSUE=Endothelial
                                                                                                                                                                                                                                                                                                             IN LEUKOCYTE-ENDOTHELIAL CELL ANDESION. INTERACTS WITH THE BETA-1
INTEGRIN VLA4 ON LEUKOCYTES, AND MEDIATES BOTH ADHESION AND SIGNAL
TRANSDUCTION. THE VCAMI/VLA4 INTERACTION MAY PLAY A
PATHOPHYSTOLOGIC ROLE BOTH IN IMMUNE RESPONSES AND IN LEUKOCYTE
EMIGRATION TO SITES OF INFLAMMATION.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN (LONG ISOFORM) OR
ATTACHED TO THE MEMBRANE BY A GPI-ANCHOR (SHORT ISOFORM)
ALTERNATIVE PRODUCTS: THE GPI-ANCHORED AND THE TRANSMEMBRANE
ISOFORMS ARE PRODUCTS: THE GPI-ANCHORED AND THE TRANSMEMBRANE
ISOFORMS ARE PRODUCTS: THE GPI-ANCHORED AND THE TRANSMEMBRANE
ISOFORMS ARE PRODUCTS: THE GPI-ANCHORED AND THE TRANSMEMBRANE
ISOFORMS ARE PRODUCTS: THE GPI-ANCHORED AND THE TRANSMEMBRANE
ISOFORMS ARE PRODUCTS: EXPRESSED ON INFLAMED VASCULAR ENDOTHELIUM, AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Immuno1.
                                                                                                                                                                                                                                                                                                      TISSUE SPECIFICITY: EXPRESS WELL AS ON MACROPHAGE-LIKE
                                                                                                                                                                                                                                                                                                 AND
                      L22352;
L22353;
L22354;
                                                            L22355;
L22301;
L22349;
L22350;
L22351;
                                                                                                                                              M84487;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of murine vascular cell adhesion
                                                                                                                                                                                                                                                                                              INFLAMED TISSUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of an inflammation-specific phosphatidyl i murine vascular cell adhesion molecule-1.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18:387-391(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AUG-1994) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kwee L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         153:4088-4098(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Lung
                                                              AAA16921.
AAA16921.
                                                                                                                              AAA40545.1;
CAA47989.1;
                                                                                        AAA16921
                       AAA16921
                                    AAA16921
                                                  AAA16921
                                                                                                        AAA16921
                                                                                                                   AAA16921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              268:8835-8841(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Liver;
9; PubMed=7523515;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Levine J.F.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GPI-ANCHORED ISOFORM)
                                                                                                                                                                                                                                                                                                        AND DENDRITIC CELL TYPES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Labow M.A.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mapping, and analysis of
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                                                                                                                                                                                                           There are no rest
                                                                                                                                                                                   http://www.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ISOFORM)
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                                                                                                                                                                                               Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gotto A.M.
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RESULT
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Best Local :
VCA1_RAT
P29534;
01-APR-1993
01-APR-1993
30-MAY-2000
                                                                                                                                                                       VARSPLIC
CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                           DOMAIN
CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                 CARBOHYD
CARBOHYD
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MGD; MGI:99926; Vcaml.
InterPro; IPR003006; -
Pfam; PF00047; 19; 5.
Immunoglobulin domain; Glycoprotein; Cell adhesion; Transmembrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL;
EMBL;
EMBL;
EMBL;
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EMBL;
                                                  RAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     GPI-anchor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PIR; JS0674; JS0674.
PIR; JN0581; JN0581.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; L12541; AAC37607.1; EMBL; U42327; AAB88576.1;
                                                                                                                                                                                                                                                  CARBOHYD
                                                                                                                                                                                                                                                                                         DOMAIN
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661 QDAGIYECESKT 672
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                                                                                             1 EDSGLYQCEAAT 12
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U12882; AAA80011.1

U12875; AAA80013.1

U12872; AAA80013.1

U12873; AAA80014.1

U12873; AAA80014.1

U12877; AAA80016.1
                                                                                                                          Similarity 50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 )8431; AAA40546.
L2884; AAA64832.
                                                                                                                                                                      346
693
739
  (Rel.
(Rel.
(Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Signal;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB60664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB60661
                                                                                                                                                                         AA;
                                        STANDARD;
25, Created)25, Last sequence update)39, Last annotation update)
                                                                                                                                                                       739
693
81317
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739
198
198
287
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575
575
675
225
227
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Alternative
                                                                                                                                    70.3%;
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IG-LIKE C2-TYPE DOMAIN 1.

IG-LIKE C2-TYPE DOMAIN 2.

IG-LIKE C2-TYPE DOMAIN 3.

IG-LIKE C2-TYPE DOMAIN 4.

IG-LIKE C2-TYPE DOMAIN 5.

IG-LIKE C2-TYPE DOMAIN 5.

IG-LIKE C2-TYPE DOMAIN 7.

N-LINKED (GLCNAC...) (POT
                                                                                                                         Score 45; DB 1; Length 739; Pred. No. 0.62; Indels
                                                                                                                                                                                         DGRMKSQIINGHQLTVHLMFAKSFYFICYLCLYLAL (IN GPI-ANCHORED ISOFORM).
MISSING (IN GPI-ANCHORED ISOFORM).
                                                                                                                                                                                                                    N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
EKPFIYDISPGSQVAAQVGDSVYLTCAAIGCDSPSF
                                                                                                                                                                                                                                                                                                                                                                                                      VASCULAR CELL ADHESION PROTEIN 1, GPI-ANCHORED ISOFORM.
                                                                                                                                                                                                                                                                                                                                                                           GPI-ANCHOR (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                             REMOVED IN MATURE GPI-ANCHORED ISOFORM
                                                                                                                                                                                                                                                                                                                                                                                                                        VASCULAR CELL ADHESION PROTEIN TRANSMEMBRANE ISOFORM.
                                                                                                                                                                                                                                                                                                                                                                 EXTRACELLULAR
                                       PRT;
                                                                                                                                                                        3D2134C341E5E449 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   splicing.
                                       739
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                                                                                                                          0
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RESULT 13
NCM2\_MOUSE
ID NCM2\_M

NCM2\_MOUSE

STANDARD;

PRT;

837 AA

D

661 QDAGVYECESKT 672

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ROCCOS
  δĀ
                                               Query Match
Best Local S
Matches 6
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SEQUENCE
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DOMAIN
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SIGNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fcentities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VASCULAR CELL ADHESION PROTEIN 1 PRECURSOR (V-CAM VCAM1 OR VCAM-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00047;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIR; JS0675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; M84488; AAA42332.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        - !- SIMILARITY: CONTAINS 7 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=92181437; PubMed=1371918;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE-Lung;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       [nterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HSSP; P19320; 1VCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mmunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.

TISSUE SPECIFICITY: EXPRESSED ON INFLAMED VASCULAR ENDOTHELIUM, AS
TISSUE SPECIFICITY: EXPRESSED ON INFLAMED VASCULAR ENDOTHELIUM, AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AND
EDSGLYQCEAAT
                                               h 68.8%;
Similarity 50.0%;
6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AS ON MACROPHAGE-LIKE AND DENDRITIC CELL TYPES INFLAMED TISSUE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; IPR003006; 00047; ig; 5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        domain; Glycoprotein; Cell adhesion; Transmembrane;
     12
                                                                                                                                                                     198
379
379
496
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675
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8124
81246
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739
698
720
739
91
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                                                                                                                                                                                                             CYTOPLASMIC (POTENTIAL).

IG-LIKE C2-TYPE DOMAIN 1.

IG-LIKE C2-TYPE DOMAIN 2.

IG-LIKE C2-TYPE DOMAIN 3.

IG-LIKE C2-TYPE DOMAIN 4.

IG-LIKE C2-TYPE DOMAIN 5.

IG-LIKE C2-TYPE DOMAIN 7.

IG-LIKE C3-TYPE DOMAIN
                                                    Ġ
                                               Score 44; DB 1;
Pred. No. 0.96;
5; Mismatches
                                                                                                                                                                                                N-LINKED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  POTENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        VASCULAR CELL
EXTRACELLULAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Rattus
                                                                                                                                                                          5C608E5A1A1B100C
                                                                                                                                                                                              (GLCNAC. .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (POTENTIAL).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADHESION PROTEIN 1
                                                    1;
                                                                                            Length 739;
                                                                                                                                                                       CRC64;
                                                    Indels
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(POTENTIAL).
(POTENTIAL).
(POTENTIAL).
(POTENTIAL).
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                                                    Gaps
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A ALENIAMS A., Bohm S.;

A Alenius M., Bohm S.;

"Identification of a novel neural cell adhesion molecule-related gene
"With a potential role in selective axonal projection.";

L J. Biol. Chem. 272:26083-26086(1997).

C -!- FUNCTION: MAY PLAY IMPORTANT, ROLES IN SELECTIVE FASCICULATION AND
C ZONE-TO-ZONE PROJECTION OF THE PRIMARY OLFACTORY AXONS.

C -!- SUBCELLULAR LOCATION: TYPE I.MEMBRANE PROTEIN (LONG FORM) AND
C -!- SUBCELLULAR LOCATION: TYPE I.MEMBRANE PROTEIN (LONG FORM) AND
C -!- SUBCELLULAR LOCATION: TYPE I.MEMBRANE PROTEIN COLFACTORY AND
C -!- TISSUE SPECIFICITY: EXPRESSED IN SUBSETS OF BOTH OLFACTORY AND
C -!- SIMILARITY: CONTAINS 5 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS.
C -!- SIMILARITY: CONTAINS 2 FIBRONECTIN TYPE III-LIKE DOMAINS.
                                                                                                                                                                                                                                                                           EMBL; AF001287; AAB69125.1; -.
EMBL; AF001286; AAB69124.1; -.
EMBL; AF016619; AAC53375.1; -.
MGD; MGI:1095738; Ocam.
InterPro; IPR001777; -.
InterPro; IPR003006; -.
 DISULFID
DISULFID
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DISULFID
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DOMAIN
                                                                                                                                                                                                                                                    Pfam; PF00041; fn3; Pfam; PF00047; ig;
                                                                                                                                                               DOMAIN
                                                                                                                                                                           TRANSMEM
                                                                                                                                                                                          DOMAIN
                                                                                                                                                                                                                  SIGNAL
                                                                                                                                                                                                                                                                                                                                                                    use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See or send an email to license@fisb-sib.ch)
                                                                                                                                                                                                                                             Cell adhesion;
                                                                                                                                                                                                                             Immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                    use
                                                                                                                                                                                                                                                                                                                                                                                                                               the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN-C57BL/6J; TISSUE-Olfactory epithelium; MEDLINE-97476194; PubMed-9334170;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. (LONG AND SHORT FORMS).
STRAIN-BALB/C: TISSUE-Olfactory neuroepithelium;
MEDLINE-97368238; PubMed-9221781;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mus musculus (Mouse)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   035136: 035962;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUL-1998 (Rel. 36, Last annotation update)
NEURAL CELL ADHESION MOLECULE 2 PRECURSOR (N-CAM 2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              axons.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neurosci. 17:5830-5842(1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   related to zone-to-zone projection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yoshihara Y., Kawasaki M., Tamada A.,
Kagamiyama H., Mori K.;
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                                                                                                                                                                                                                            domain;
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                                                                        CYTOPLASMIC (POTENTIAL).
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C3-TYPE-DOMAIN.
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                                                                                                                                                                                                               POTENTIAL
                                                                                                                                                                                                                   Glycoprotein; Repeat;
; GPI-anchor; Alternative splicing.
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Sciurognathi;
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n of olfactory
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thi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                          as its content
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Murinae; Mus
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NCM2_HUMAN
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Genomics 43:43-51(1997)
-1- FUNCTION: MAY PLAY IMPORTANT ROLES IN SELECTIVE PASCICULATION
ZONE-TO-ZONE PROJECTION OF THE PRIMARY OLFACTORY AXONS.
-1- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.
-1- SUBCELLULAR LOCATION: EXPRESSED MOST STRONGLY IN ADULT AND FETAL
                                Pfam; PF00041; fn3; 2. Pfam; PF00047; ig; 5:
                                                       InterPro; IPR001777; -.
InterPro; IPR003006; -.
                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCM2_HUMAN
015394;
15-JUL-1998
                         Cell adhesion;
                                                                                MIM; 602040;
                                                                                           EMBL;
                                                                                                            entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                    use by non-profit institutions as long modified and this statement is not removed.
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Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
             mmunoglobulin
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Paoloni-Giacobino
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CARBOHYD
VARSPLIC
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PCLoning of a novel human neural cell adhesion molecule gene (NCAM2)
that maps to chromosome region 21q21 and is potentially involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                    NCAM2 OR NCAM21
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15-JUL-1998 (Rel. 36, Last annotation update)
NEURAL CELL ADHESION MOLECULE 2 PRECURSOR (N-CAM
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domain;
        Transmembrane; Glycoprotein; Repeat; domain; Signal.
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2 FIBRONECTIN TYPE III-LIKE DOMAINS
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ITRRMCGKKSGSSGKSKELEEGKAAYLKDGSKEPIVEMRTE
DERITMHEDGSPVNEPNETTPLTEPEKLPLKEENGKEVLNA
ETIEIKVSNDIIQSKEDDIKA -> NCCEANKGENGGQSWH
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01-CCT-2000 (Rel
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SEQUENCE
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01-OCT-2000 (Rel. 40, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPTOR 2 PRECURSOR (EC (VEGFR-2) (KINASE INSERT DOMAIN RECEPTOR) (FLK-1).
KDR OR FLK1.
                Oncogene [4]
                                                              TISSUE-Umbilical vein;
MEDLINE-92019839; PubMed-1656371;
                                                                                                        TISSUE-Umbilical
Yu Y., Whitney R
Coding region for
                                                                                                                                                 Submitted
                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
SEQUENCE OF 1-22 I
MEDLINE=96032749;
                                                        Terman B.I.,
                                                                                                 Submitted
                                                                                                                                                Yin L.Y., Wu Y., Patterson C.;
"Full length human KDR/flk-1 sequence.";
Submitted (DEC-1997) to the EMBL/GenBank
                                   "Identification of a new endothelial cell growth factor recepton
                                                                               SEQUENCE OF 3-1356 FROM N.A.
                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                         SEQUENCE FROM N.A
                                                                                                                                                                                         NCBI_TaxID=9606
                                                                                                                                                                                                                 iomo sapiens (Human).
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EDAGIYRCQA
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                                                  T.B.;
                                                                                                                                                                                                                                                                                                                                                           Similarity 6; Conser
                        kinase.";
6:1677-1683(1991)
                                                                                               itney R.G., Sato J.
egion for human VEG
(MAY-1998) to the
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232
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                                                         Carrion M.E.,
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FROM N.A.; PubMed=7559454;
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Primates;
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                                                         Kovacs
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE-III.
FIBRONECTIN TYPE-III.
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receptor J. Biol. [5]
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Harber E.;
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Biochem. Biophys. Res. Commun. 187:1579-1586(1992).
Biochem. Biophys. Res. Commun. 187:1579-1586(1992).
-I- FUNCTION: RECEPTOR FOR VEGF-OR VEGF-C. HAS A TYROSINE-PROTEIN KINASE ACTIVITY. THE VEGF-KINASE LIGAND/RECEPTOR SIGNALING SYSPLAYS A KEY ROLE IN VASCULAR DEVELOPMENT AND REGULATION OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Harber E.;
"Cloning and functional analysis of the receptor for vascular endothelial growtl receptor for vascular 11-23118(1995).
                                                                                                                  This
                                                                                                                                                                                                                                                                                                                                                        MEDLINE=93038639; PubMed=1417831;
Terman B.I., Dougher-Vermazen M., Carrion M.E., Dimitrov
Armellino D.C., Gospodarowicz D., Boehlen P.;
"Identification of the KDR tyrosine kinase as a receptor
              modified and this statement is not rementities requires a license agreement
                                                                        the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION
                                                                                               between
send
                                                                                                                                                                          SUBCELLULAR LOCATION: TYPE SIMILARITY: BELONGS TO THE PROTEIN KINASES.
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                                                                                                                                                                                                                                          VASCULAR PERMEABILITY
                                                                                               SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the \Xi
  an email to license@isb-sib.ch).
                                                        non-profit institutions as long
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                      is not removed. Usage hadresmore
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CSF-1/PDGF
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                  (See http://www.isb-sib
                                                                            There are no
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DR PROSITE; PS00107; PROTEIN\_KINASE\_ATP; FALSE\_NEG.
DR PROSITE; PS00109; PROTEIN\_KINASE\_TYR; 1.
DR PROSITE; PS00240; RECEPTOR\_TYR\_KIN\_II; 1.
DR PROSITE; PS00240; RECEPTOR\_TYR\_KIN\_II; 1.
DR PROSITE; PS50011; PROTEIN\_KINASE\_DOM; 1.
W Signal; Transferase; Tyrosine-protein kinase: Protein Glycoprotein; Phosphorylation: Numerical Signal. CARBOHYD CARBOHYD CARBOHYD NP\_BIND BINDING ACT\_SITE TRANSMEM DOMAIN DOMAIN MIM; 191306; InterPro; IP CARBOHYD CARBOHYD EMBL; EMBL; EMBL; EMBL; HSSP; DOMAIN DOMAIN DOMAIN DOMAIN InterPro;
InterPro; CHAIN DOMAIN DOMAIN DOMAIN DOMAIN AF035121; AAB88005.1; -. AF063658; AAC16450.1; -. x89776; L04947; x61656; IPR001245; IPR001824; PR000719; 765 790 790 141 141 345 CAA43837:1; AAA59459:1; 1356 CYTOPLASMIC (POTENTIA IG-LIKE DOMAIN 1 ... IG-LIKE DOMAIN 2 ... IG-LIKE DOMAIN 3 ... IG-LIKE DOMAIN 4 ... IG-LIKE DOMAIN 5 ... IG-LIKE DOMAIN 6 ... IG-LIKE DOMAIN 7 ... PROTEIN KINASE ... ATP (BY SIMILARITY) ... ATP (BY SIMILARITY) ... ATP (BY SIMILARITY) ... BY SIMILARITY N-LINKED (GLCNAC ... N-LINKED (GLCNAC ... INTERD GLCNAC ... IG-LIKED (GLCNAC ... INTERD GLCNAC ... INTERD (GLCNAC ... INTERD GLCNAC .. N-LINKED N-LINKED VASCULAR RECEPTOR POTENTIAL EXTRACELLULAR (POTENTIAL) (GLCNAC. (GLCNAC. (GLCNAC. (GLCNAC ENDOTHELIAL GROWTH (POTENTIAL). Immunoglobulin Receptor; (POTENTIAL).
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Search completed: June
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Best Local Similarity 58.3%;
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13.062 Million cell updates/sec
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Q90784 ginglymosto
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Q18100 caenorhabdi
Q63155 rattus norv
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o sapien	caenorhabdi	ginglymosto	gallus gal	aedes aegyp	nglymc	homo sapien	xenopus lae	hirudo medi	rattus norv	rattus norv	s musculu	s musculu	s musculu	drosophila	taurus	cavia porce	cavia porce	ginglymos	drosophila	drosophila	xenopus lae	brachydanio	s musculu	s musculu

## ALIGNMENTS

RESULT Q92663

RABBE

Q92663 Q92663;

PRELIMINARY;

PRT;

374 AA.

01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation update)

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RESULT 2 Q92495 ID Q92495 PRELIMINARY; PRT; 375 AA.	Qy 1 EDSGLYOCEAAT 12 	Query Match  89.1%; Score 57; DB 4; Length 374;  Best Local Similarity 91.7%; Pred. No. 0.0032;  Matches 11; Conservative 0; Mismatches 1; Indels 0; Gaps 0;	SQ SEQUENCE 3/4 AA; 42032 MW; D3DD3398CEEA099 CKC04;	PRODOM; PD002534; -; 1.	DR INTERPRO; IPRO03006;	KE J. CIIN. INVEST. 90:2102-2109(1992).  DR EMBL; L03418; AAA36049.1;	RT cells,";	RT "Novel Fc gamma receptor I family gene products in human mononuclear	Kimberly R.P.;	RX MEDIINE-93055454; PubMed-1430234; C. Ralmon T. F. Ralm	RC TISSUE=BLOOD;	RP SEQUENCE FROM N.A.	 Primates;	OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	OS Al. OS Homo sapiens (Human).	DE FC GAMMA RECEPTOR I	
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Q90284;
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01-NOV-1996 (Tremblrel. 01, L
01-MAY-2000 (Tremblrel. 13, L
   Giordano S., Laessing U., Lot
Submitted (APR-1996) to the E
EMEL; U55211; AAA99159.1; --
HSSP; P20241; 1CFB.
INTERPRO; IPR001777; --
INTERPRO; IPR003006; --
                                                                                                                                                               Carassius auratus (Goldfish).
Eukaryota; Metazoa; Chordata; Craniata;
Actinopterygii; Neopterygii; Teleostei;
Cypriniformes; Cyprinidae; Cyprininae; (
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01-FEB-1997.
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01-OCT-2000
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                                                                                                                SEQUENCE FROM N.A.
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SEQUENCE
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Submitted (DEC-1992)
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Benech P.D.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          . Exp. Med. 176:1115-112
MBL; M91555; AAA58414.1;
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-FEB-1997 (TremBLrel. 02, Last sequence update)
-CCT-2000 (TremBLrel. 15, Last annotation update)
-CAMMA RECEPTOR TYPE I (FC GAMMA RIB=FC GAMMA RECEPTOR).
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S45704;
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M91551; AAA58414.1; JOINED
M91552; AAA58414.1; JOINED
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1PR001777; -.
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4; AAA58414.1;
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s (Human)
chordata; (
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PubMed=1402657;
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he EMBL/GenBank/DDBJ
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Last annotation updat
ULE ANTIGEN E587.
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Euteleostei;
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0.0032;
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                                                                                          Stuermer C.A.O.;
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; Ostariophysi;
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Q90874;
Q1-NOV-1996
Q1-NOV-1996
Q1-OCT-2000
                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N:A.
SEQUENCE FROM N:A.
MEDLINE=99045686; PubMed=9826702;
Diaz M., Greenberg A.S., Flajnik M.F.;
"Somatic hypermutation of the new antigen receptor gene (NAR) nurse shark does not generate the repertoire: possible role ir antigen-driven reactions in the absence of germinal centers."

antigen-driven reactions in the absence of germinal centers."

Na+1 Acad. Sci. U.S.A. 95:14343-14348(1998).
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PFAM; PF00047; ig
PRINTS; PR00014;
SEQUENCE 1232 A
                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute. Eukaryota; Metazoa; Neoqnathae; Galliformes; Phasianidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q9YHT0;
01-MAY-1999
01-MAY-1999
01-MAY-2000
SEQUENCE FROM N.A.
MEDLINE=92175529; PubMed=1531799;
Guida S. Heguy A. Welli M.;
"The chicken IL-1 receptor: differential
                                                                                                    IL-1
                                                                                                                                                                                                                                                                                                      NON_TER
                                                                                                                                                                                                                                                                                           SEQUENCE
                                                  NCBI_TaxID=9031;
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Eukaryota; Metazoa; Chordata; Craniata; V
Elasmobranchii; Galeomorphii; Galeoidea;
Ginglymostomatidae; Ginglymostoma.
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105 AA;
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(TrEMBLrel. 10,
(TrEMBLrel. 13,
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4; FNTYPBIII.
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01-NOV-1998
01-MAY-2000
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CDO.
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Q63198;
01-NOV-1996
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EMBL; D38492; BAA07504.1;'-.
HSSP; P40189; 1BQU.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Takenawa T.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-2000 (TremBLrel. 13, Last annotation Neural ADHESION MOLECULE F3, COMPLETE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-AUG-1998 (TrEMBLrel. 01-MAY-2000 (TrEMBLrel.
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HSSP; P14778; 1ITB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rat brain.
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MEDLINE=95295987; PubMed=7777204;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10116;
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e 111:239-243(1992).
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EDEGLYECEA 312
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8; Conserv
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8; Conservative
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00047; ig; 6.
PS00092; N6_MTASE; UNKNOWN_1.
1021 AA; 113494 MW; FCBDC13055EE5C68 CRC64;
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                                            (TrEMBLrel. 05, (TrEMBLrel. 08, (TrEMBLrel. 13,
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63995 MW;
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1; Mismatches
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Last annotation update)
                                                                                      Created)
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Pred. No. 0.5;
1; Mismatches
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Sciurognathi; Muridae;
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4DB56DD7B1BB17AA CRC64;
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COMPLETE CDS.
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No. 0.28;
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; Murinae; Rat
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PFAM; PF00041; fn3; 3.
PFAM; PF00047; ig; 5.
SEQUENCE 1256 AA; 1
                                                                                    Signal.
SIGNAL
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                                                                                                                                                                                                                                                                                                                                              Q63611;
01-NOV-1996
                                                             SEQUENCE
                                                                          CHAIN
                                                                                                                                               J. Biol. Chem. 269:15601-
                                                                                                                                                                                               STRAIN=BDIX; TISSUE=COLON TUMOR; MEDLINE=94253144; PubMed=8195207;
                                                                                                                                                                                                                                                          Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae; Rat
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Submitted (S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "CDO: an oncogene-, serum-, and anchorage-regulated member of the Ig/fibronectin type III repeat family.";
J. Cell Biol. 138:203-213(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. MEDLINE=97362072; | Kang J.S., Gao M.,
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
                                                                                                            PFAM; PF00047; ig;
                                                                                                                                                                                  Chadeneau C., LeMoullac B., Denis M.G
                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                              NCBI_TaxID=10116;
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                                                                                                                         INTERPRO;
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 Similarity 7; Conserv
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9; Conser
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(SEP-1998) to
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412
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                                                                                                                                                269:15601-15605(1994).
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, Feinleib J.L.,
                                                             44509
            73.4%;
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EMBL/GenBank/DDBJ
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EMBL/GenBank/DDBJ
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           Score 47;
Pred. No.
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TUMOR-ASSOCIATED ANTIGEN.
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1.32;
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                      Length 412;
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  Indels
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; Murinae; Rattus
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Best Local
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Q60977;
Q1-NOV-1996
        EMBL; U35836; AAB17502.1;
HSSP; Q13740; 1KJC.
MGD; MGI:107741; Taal.
INTERPRO; IPR001064; -.
                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                  TUMOR-ASSOCIATED GLYCOPROTEIN
                                                                                                                                                                                            01-OCT-2000
                                                                                   Chadeneau C., LeMoullac B., LeCabellec
                                                                                                          STRAIN=BALB/C
                                                                                                                 SEQUENCE FROM: N.A.
                                                                                                                                                                 Mus musculus (Mouse)
                                                                                                                                                                          TAA1 OR TAGE4.
                                                                                                                                                                                                       01-NOV-1996
                                              mmunoglobulin superfamily.";
famm. Genome 7:636-637(1996).
                                                                              enis M.G.
                                                                                              MEDLINE=96327659; PubMed=8678999;
                                                                                                                                    CBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted
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"Structure of the rat
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NCBI_TaxID=10116;
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                                                                Isolation and chromosomal location
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EDEGIYECQIAT 134
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IPR003006; -
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                                                                                                                                           Chordata;
Rodentia;
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Rodentia;
                                                                                                                                                                                                                                                                                                                                                                  44483 MW;
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EMBL/GenBank/DDBJ databases
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Pred. No.
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                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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INTERPRO; IPR003006; -.
PFAM; PF00041; fn3; 6.
PFAM; PF00047; -19; 6.
SEQUENCE 1287 AA; 144
Q63155 PRELIMINARY;
Q63155;
01-NOV-1996 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                      STRAIN-BRISTOL N2;
Du Z., Gattung S.;
"The sequence of C. e
Submitted (MAR-1996)
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01-MAY-2000
01-OCT-2000
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STRAIN=BRISTOL N2;
MEDLINE=99069613; PubMed=9851916;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea; Rhabditidae; Peloderinae; Caenorhabditis.
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SEQUENCE
                                                                                 168
                                                                                                                                                                                                                                                                                STRAIN-BRISTOL N2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     investigating biology. The C
Science 282:2012-2018(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genome sequence of the nematode C.
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                                                                                                     2 DSGLYQCEA 10
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8; Conserv
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415 AA; 45280 MW; 159AF8D60EF90D0F CRC64;
                                                                                                                                                                                                                                                                                                                 (FEB-1996)
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EMBL/GenBank/DDBJ databases.
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Pred. No. 1.1;
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01-OCT-2000
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       Bouchon A., Dietrich J., Colonna M.;

Bouchon A., Dietrich J., Colonna M.;

"Cutting edge: inflammatory responses can Be triggered by TREM-1, novel receptor expressed on neutrophils and monocytes.";

J. Immunol. 164:4991-4995(2000)

EMBL; AF287008; AAF90197.1; -.

EMBL; AF196329; AAF71694.1; -.
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Culotti J.G., Tessier-Lavigne
"Deleted in Colorectal Cancer
Cell 87:175-185(1996).
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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EMBL; M32291; AAA41086.1;
HSSP; P56276; ITLK
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MEDLINE=90100559; PubMed-2294591;
Fearon E.R., Cho K.R., Nigro J.M.
Ruppert J.M., Hamilton S.R., Prei:
                                                                                                                                                       SEQUENCE FROM N.A. MEDLINE-20261626; PubMed-10799849;
                                                                                                                                                                                                                                                                        Gingras M.C., Lapillonne H. Monocytic differentiation
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Mammalia; Eutheria;
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Metazoa; Chordata; C
Metazoa; Primates; (
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Thomas
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; Murinae; Rattus
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G., Kinzler
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Q28106;
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               Warren J.T.
Kuwada J.Y.;
                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; l
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Cypriniformes; Cyprinidae; Rasborinae; Danio.
                                                                                                                                                                                             Brachydanio rerio (Zebrafish) (Zebra danio)
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01-MAY-2000
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Mammalia; Eutheria;
Bovidae; Bovinae; B
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TE; PS00092; N6_MTASE;
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                                      Kanki J.P.,
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'Molecular

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and developmental

expression

analysis of the

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd

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Listing first 45 summaries
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Human FCRI (CDNA C
Human macrophage-s
Human macrophage-s
Human FCRI (CDNA C
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Fc receptor
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Arabidopsis thalia			G23166	. 21	156	57.4	်. ယ	
ted			208	21	89	57.4	ω	
			531	21	586		36	
sezia func			2977	18	507	•	36.	
Human secreted pro			474	21	359	59.0	36	
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PR0329			4169	20	359	•	36	
ระ			Y94336	21	358	•	36	
bidopsis			G42149	21	321	•	36	
gene produc			W14450	17	288	•	36	
S. pneumoniae der			Y86103	19	132		36	
			G53158	21	753	60.7	37	
is thal			G53159	21	715	•	3.7	
S			G52083	21	673	60.7	37	
Arabidopsis thali			G52084	21	637	60.7	37	
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idopsis thal			G13324	21	523	62.3	38	
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Arabidopsis thal			G13326	21	431	•	38	
idopsis			G28824	21	421	•	42	
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ŝ	<i>)</i>	•	G12849	21	163	•	42	
idopsi			G12850	21	149	•	42	
Hybrid Fc(gamma)R			R12428	12	410	•	50	
Human Fc receptor			W86195	20	377	•	52	
Human macrophage-			Y96134	21	374	•	53	
Human macrophage-	*		618	21	344	•	53	
Human cancer asso			B43683	21	399	100.0	61	
Human high affini			9622	21	374	8	61	

## ALIGNMENTS

RESULT W60559

W60559;

18-AUG-1998

(first entry)

W60559 standard; peptide; 12

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Oligopeptide from extracellular domain of CD64.

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies Synthetic. cell transformation; mutation; anti; oncogene. WPI; 1998-240964/21. De Kruif CA, 08-OCT-1996; 07-OCT-1997; 16-APR-1998 WO9815833-A1. Extracellular domain; CD64; identification; antibody; (UYUT-) RIJKSUNIV UTRECHT. mmunohistochemical; Logtenberg T; 96EP-0202791 97WO-NL00557 immunofluorescent analysis; detection,

Synthetic oligor domain of CD64.

oligopeptides W60537-61

Example 1; Page 29;

40pp; English

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Best Local Similarity
Matches 12; Conserv
                                                                                                                  Baell JB,
Maxwell K
                                                                                                                                                                                                     11-SEP-1998;
06-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                               For receptor; Fc-gammaRIIa; human; FCR; model; three-dimension; 3-D; atomic coordinate; bioactive compound design; computer-assisted; drug design; therapy; inhibitor; Fc-gammaR; Fc-epsilonR; IgG; IgE; tissue damage; hypersensitivity; inflammatory cell recruitment; inflammatory modulator; Fc-gammaRIa; immune function regulation;
                                                                                                                                                                                                                                                       04-FEB-1999;
                                                                                                                                                                    (ILEX-) ILEXUS PTY LTD.
                                                                                                                                                                                                                                                                                                                          WO9940117-A1
                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                          anti-inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to epitopes on the same monometric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or
                                                                                                                                                                                                                                                                                       12-AUG-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y33183 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic oligopeptides W60537-61 are derived from the extracellular domain of CD64. They were synthesised on 25 polyethylene rods as adjacent, non-overlapping sets. Peptide specific phages were isolated by affinity binding to the CD64-covered rods. The specification describes a method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a renlicable displaying the peptide on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  by mutation in (anti)oncogenes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antibodies. Selected antibodies are used, e.g. for immunohistocheminmunofluorescent analysis, and also to detect cell transformation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 edgnvlkrspel 12
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ilarity 100.0%;
Conservative (
                                                                                                                 pa V, Garrett TPJ,
McCarthy TD, McKe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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98US-0073972.
                                                                                                                                                                                                                                                     99WO-IB00367.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein fragment.
                                                                                                                                                                                                                                                                                                                                                                                      immunoprotective; src-gammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              261 AA.
                                                                                                               TPJ, Hogarth McKenzie IFC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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                                                                                                             PM, Matthews
Pietersz GA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 12;
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Three-dimensional structures and models of Fc receptors, useful in computer-assisted drug design  $% \left( 1\right) =\left\{ 1\right\} =\left$ 

This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residu while clone p98/X2 predicts a Thr residue. At position 58, p90 '

residue

Example 10;

Page 94a;

160pp; English.

WPI; 1999-539978/45

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PDX
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Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                       Seed
                                                                                                               New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                     WPI; 1992-056864/07.
N-PSDB; Q21179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reduce IgG- or IgE-mediated tissue damage, e.g. IgG- or IgE-mediated hypersensitivity, recruitment of inflammatory cells or release of inflammatory modulators. The therapeutic compositions can also be used to treat disease in animals. The knowledge of the 3-D structure of Fc-gamma compounds that regulate immune function and inflammation in an animal, including humans (i.e. structure based drug design). For example, chemical compounds can be designed to block binding of immunoglobulin to an Fc receptor protein using various computer programs and models. The products of the invention have anti-inflammatory and immunoprotective activity. This sequence represents the human sFC-gammaRI protein fragment described in the method of the invention.
                                                                                                                                                                                                                                                       (GEHO-) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                          13-JUL-1990;
                                                                                                                                                                                                                                                                                                                        15-JUL-1990; '
                                                                                                                                                                                                                                                                                                                                                            23-JAN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rapid immunoselection cloning technique; cell surface antigen, immunodiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage-specific FcRI receptor encoded by clone p98.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R22549 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                              WO9201049-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotides, peptides, peptidomimetics and small organic molecules by computer-assisted methods of drug design. Therapeutic compositions that inhibit the activity of Fc-gammaR or Fc-epsilonR can be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 261 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the FCR can be used to design bloactive chemical compounds,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                representing a three-dimensional (3-D) structure that substan conforms to the specified atomic coordinates. Computer model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes a novel model of an Fc receptor (FCR) protein representing a three-dimensional (3-D) structure that substantially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 11; Page 316-317; 326pp; English
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                                                                                                                                                                                                                     В,
                                                                                                                                                                                                                     Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100:0%; Score 61; Dilarity 100:0%; Pred. No. 0.0 Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                      90US-0553759
                                                                                                                                                                                                                                                                                                                        90WO-US04986
                                                                                                                                                                                                                     Amiot M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.001;
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                                                                                                                                 it - for
                                                                                                               neoplasms,
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                                                                                                               etc.
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RESULT
R91439
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Best Local
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          The amino acid sequence (R91439) of human FCRI was detd. from a CDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T14719), macrophages.
                                                                                                                                                                                                                                                                                                                                                                                                                                   01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
                                                                                                                                                                                          and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning doma.
                                                                                                                                                                                                                                                         WPI; 1996-200279/20
N-PSDB; T14718.
                                                                                                                                                                                                                                                                                                                   Aruffo
                                                                                                                                                                                                                                                                                                                                                                                              13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cell surface antigen; cloning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human FCRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R91439 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                                                       Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     portion
                                                                                                                                                                                                                                                                                                                                                       (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ortion contains six potential N-linked glycosylation sites and six ys residues distributed among three C2 set 1g-related domains. fusion protein of FCRI and a receptor ligand will be helpful to ncrease the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 EDGNVLKRSPEL 12
the p98 product
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               a short, highly charged cytoplasmic domain. The extracellular tion contains six potential N-linked glycosylation sites and si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     l Similarity
12; Conserv
                                                                                                                                                                                                                                                                                                                   P,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                   92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                              900S-0553759
930S-0139273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88US-0160416
                                                                                                                                                                                                                                                                                                                 В;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 61;
100.0%; Pred. No.
tive 0; Mismatc
  truncated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "amino acid 58 is Leu in clone translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunoselection; immunotherapy;
  compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.0014;
with those of the pl35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13;
                                                                                                                                                                                                                  useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        domain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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Query Match
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Matches 12; Conser

Conservative

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100.0%;

Score 61; Pred. No. Mismatches

DB 13;

Length 374;

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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            . Sequence
                                                                                                                                                          This amino acid sequence was predicted from the cDNA sequence of cDNA clone pl35. It differs from the sequence predicted from clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; pl35 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, pl35 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and p90 products (see also R91438 and of FCRI and a receptor ligand will be potency of antibodies in therapy.
                            distributed among three C2 set Ig-related domains of FCRI and a receptor ligand will be helpful to i potency of antibodies in therapy.
                                                                                             hydrophobic membrane-spanning domain, and a short, highly cytoplasmic domain. The extracellular portion contains six
                                                                                                                            typical features of a type I integral membrane protein and a short hydrophobic signal sequence, a single 21-residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R20811;
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                                                                                                                                                                                                                                                                                                                                                                                                          Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rapid immunoselection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human macrophage-specific FcRI receptor encoded by clone p135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-1992
                                                                                potential N-linked glycosylation sites and six Cys residues
                                                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                        New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                         N-PSDB; Q21178
                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-056864/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                          (GEHO-) GEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JAN-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ch 100.0%; l Similarity 100.0%; 12; Conservative (
                                                                                                                                                                                                                                                                            10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            344 AA;
                                                                                                                                                                                                                                                                         Page
                                                                                                                                                                                                                                                                                                                                                                                                                                          HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90US-0553759
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                                                                                                                                                                                                                                                                          94a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cloning technique;
affinity receptor.
                                                                                                                                                                                                                                                                          160pp;
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                                                                                                                                                                                                                                                                          English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell surface antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . A fusion protein in increasing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                            neoplasms,
                                                                                                                                                                                                                                                                                                                             it -
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                                                                 A fusion
                                                                                                             highly charged
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                                                                                                                                                                                                                                                                                                            etc.
                                                                                                                                              include
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RESULT
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                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                        cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue and clone p90 predicts a Thr residue. At position 58, p90 predicts val while clones p135 and p88/x2 (see Q21179) both predict Leu. Sequences predicted from all 3 clones show the typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six Cys residues distributed among three C2 set Ig-related domains. A fusion protein of FCRI and a receptor ligand will be helpful to increase the notenow of antibodies in thereny
   30-OCT-1996
                                        R91438;
                                                                      R91438 standard;
                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This amino acid sequence was predicted from the cDNA sequence cDNA clone p90. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser res
                                                                                                                                                                 265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New CD53 cell surface antigen and DNA encoding it - for immuno-therapy and diagnosis of haematopoietic neoplasms,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1992-056864/07
N-PSDB; Q21180.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human macrophage-specific FcRI receptor encoded by clone p90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R22550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R22550 standard; Protein; 374
                                                                                                                                                                                              1 EDGNVLKRSPEL 12
                                                                                                                                                         edgnvlkrspel 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aruffo A,
                                                                                                                                                                                                                                                                                                                       374 AA;
                                                                                                                                                                                                                                100.0%;
ilarity 100.0%;
Conservative
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90US-0553759
                                                                    Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90WO-US04986.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amiot M;
                                                                  374 AA.
                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                              Score 61; DB 13;
Pred. No. 0.0016;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                      Of.
                                                                                                                                                                                                                                                                                                                                                      antibodies
                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                 in therapy.
                                                                                                                                                                                                                                                            Length 374;
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                                                                                                                                                                                                                            0;
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       Query Match
Best Local Similarity 100.
                                                                                                                                                   cDNA clone, p135 (T14717), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FCRI sequence, and a third clone, p98 (T14718), coded for an error (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                           The amino acid sequence (R91438) of human FCRI was CDNA clone, p135 (T14717), obtd. from a cDNA librar
                                                                                                                                                                                                                                                                                                                                                                            Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                           Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-200279/20
N-PSDB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-FEB-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cell surface antigen; cloning; immunoselection; immunotherapy; therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human FcRI (cDNA clone pl35 product)
                                                                                                                              374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9305-0139273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      89US-0379076
90US-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           88US-016041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9208-0983647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "amino acid 58 is Val in p90 clone
    translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "amino acid
translated
                                                  100.0%;
                                0
                                                  Score 61;
Pred. No.
                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25 is Thr
                                                  0.0016;
                                                                                                                                                                                                                                                                                     an FcRI was detd. from cDNA library using an
                                                                   DB 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in clone p90 and
                                                                 Length 374;
                                Indels
                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              p98
                             Gaps
                                                                                                                                                                                                             p98
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RESULT
W00859
                                                                                              В
          Human FcRI (cDNA clone p90 product).
                            30-OCT-1996
                                           W00859;
                                                       W00859 standard; Protein;
                                                                                           1 EDGNVLKRSPEL 12
|||||||||||
| 265 edgnvlkrspel 276
                                                                        8
                           (first entry)
                                                        374 AA
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Cell surface antigen; cloning; immunoselection; immunotherapy;

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F F F F F F F
                                                                         FFFXOXXXXXXX
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                immunoselection cloning method. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FcRI sequence. A fusion protein of FcRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-200279/20
N-PSDB; T14719.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aruffo A,
            Misc-difference
                                                            Misc-difference
                                                                                  Misc-difference
                                                                                                                                                 cloning.
                                                                                                                                                                                                             07-JUN-1999
                                                                                                                                                                                                                                     W80448;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-APR-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5506126-A.
                                     Misc-difference
                                                                                                                       Homo sapiens
                                                                                                                                                         Fc receptor I;
                                                                                                                                                                                   Human Fc receptor I.
                                                                                                                                                                                                                                                          W80448 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               therapy; diagnosis; vector; FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                         265 edgnvlkrspel 276
                                                                                                                                                                                                                                                                                                                                   1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                      374
                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92US-0983647.
88US-0160416.
89US-0379076.
99US-0553759.
93US-0139273.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Ã,
                                                                                                                                                           FCRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88US-0160416
                                                             /note=
23
                                                                                              Location/Qualifiers
/note= "encoded by CTG"
                       'note=
                                               'note=
                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
                                                                                                                                                         cell surface antigen; lymphocyte;
                        "encoded by GAC."
                                                                      "encoded by TGG"
                                            "encoded by CTC"
                                                                                                                                                                                                                                                                                                                                                                        Ö,
                                                                                                                                                                                                                                                                                                                                                                                    Score 61;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                  DB 17;
                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                              Length 374;
                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                        Gaps
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antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-DEC-1992;
25-FEB-1988;
                                                                                                                                                 transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                                                                                                                              adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and
                                                                                                                                                                                                                                                  expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by
                                                                                                                                                                                                                                                                                            were also obtained.
                                                                                                                                                                                                                                                                                                                                                                   This is the amino acid sequence of human Fc receptor I (FCRI), deduced from cDNA clone pl35 (see V63456) isolated using a rapimunoselection cloning method from a cDNA library expressed in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-609251/51.
N-PSDB; V63456.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US5830731-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
  animals,
                                                                                                                                                                                                                                                                                                                               from a single patient undergoing extracorporeal
                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAY-1997
                                    mmunotherapeutic applications,
                                                                                                                                                                                                                                                                                                                                                     The cDNA library was constructed from polyA RNA of cells
  including
                      of immune-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
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256 ·
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171
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103
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85
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77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note=
humans
                                                                                                                                                                                                                                                                                                               Clones p90
                                                                                                                                                                                                                                                                                            A novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "encoded by
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                                                                                                                                                                                                                                                                                            method for cloning cDNAs
                      infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ьy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bу
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              bу
                                      including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGG'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GTC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATG'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAC"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT"
                                                                                                                                                                                                                                                                                                               p98 (see x07373)
                                                                                                                                                                                                                                                                                                                                   interleukin-
                                                                                                                                                                                                                                                                                            from mammalian
                                                                                                                                                                                                                                                                                                                                                                                   (rCRI), as
a rapid
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RESULT 10
W97833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100 Matches 12; Conservative
 New cloning vector and polylinker - based on existing sequences
                           WPI; 1998-609251/51
N-PSDB; X07372.
                                                     Aruffo A,
                                                                                         13-JUL-1990;
21-MAY-1997;
                                                                     (GEHO ) GEN HOSPITAL
                                                                                                                             01-DEC-1992;
                                                                                                                                                21-MAY-1997;
                                                                                                                                                                 03-NOV-1998
                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                   US5830731-A
                                                                                                                                                                                                                                Misc-difference
                                                                                                          3-JUL-1989,
                                                                                                                     25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fc receptor I; FcRI; cell surface antigen; lymphocyte;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W97833;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W97833 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human Fc receptor I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 EDGNVLKRSPEL 12
|||||||||||
| 265 edgnvlkrspel 276
                                                    Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                8905-0379076
9005-0553759
                                                                                       90US-0553759
97US-0861205
                                                                                                                 92US-0983647.
88US-0160416.
                                                                                                                                              97us-0861205
                                                                                                                                                                                                                                                                    159
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256
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176
                                                                                                                                                                                                                                                  /note=
.171
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99
                                                                                                                                                                                                                                                                                                                                            /note=
85
                                                                                                                                                                                                                                                                                                        /note= "encoded by CAA"
                                                                                                                                                                                                                                                                                                                                                                /note=.
77
                                                                                                                                                                                                                                                                                                                                                                                60
                                                                                                                                                                                                                                                                                            /note= "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                            note=
                                                                                                                                                                                                                                                                                                                                                                                        note=
                                                                                                                                                                                                                                                                                                                                                                                                          'note=
                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
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                                                                     CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                       "encoded
                                                                                                                                                                                                   "encoded by
                                                                                                                                                                                                                    "encoded by
                                                                                                                                                                                                                                                        "encoded by
                                                                                                                                                                                                                                                                         "encoded by GGC
                                                                                                                                                                                                                                                                                                                                                                                                        "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                        "encoded by CTC"
                                                                                                                                                                                                                                                                                                                                 "encoded
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                                                                                                                                                                                                                                                                                                                                 by TCC
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                                                                                                                                                                                                                                                                                                                                                   by AAT"
                                                                                                                                                                                                                     GTC"
                                                                                                                                                                                                                                                          AAC"
                                                                                                                                                                                                    GGG"
                                                                                                                                                                                                                                       ATG"
                                                                                                                                                                                                                                                                                              CCC
                                                                                                                                                                                                                                                                                                                                                                                      CTG"
                                                                                                                                                                                                                                                                                                                                                                                                         GAC"
                                                                                                                                                                                                                                                                                                                                                                                                                                           TGG"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
   for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0:
 RESULT
W97834
                                                                                                                                                                                                                                                                                                                             g,
                                                                                                                                                                                                                                                                                                                                            Qy
                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 12; Conser
Misc-difference
                 Misc-difference
                                    Misc-difference
                                                     Misc-difference
                                                                       Misc-difference
                                                                                         Misc-difference
                                                                                                                             Misc-difference
                                                                                                                                                                                                                       Human Fc receptor
                                                                                                                                                                                                                                          07-JUN-1999
                                                                                                          Misc-difference
                                                                                                                                                Misc-difference
                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                              Fc receptor I; cloning.
                                                                                                                                                                                                                                                              W97834;
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                            265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    were also obtained.
                                                                                                                                                                                                                                                                                                                         11
                                                                                                                                                                                                                                                                                                                                                                                                                            including humans.
                                                                                                                                                                                                                                                                                                                                                                                                            374 AA;
                                                                                                                                                                                                                                                                                                                                                              100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                         (first entry)
                                                                                                                           /note=
. 23
/note=
103
                                    /note=
85
                                                                                                                                                                                                      FCRI;
                                                                       /note=
                                                                                                          /note=
                   99
                                                       /note=
77
                                                                                           45
                           'note=
                                                                                                   /note=
                                                                                                                                                                                                                                                                                                                           276
                                                                                                                                                                                                                                                                                                                                            12
      "encoded by CAA"
                                                            "encoded by
                         "encoded by TCC"
                                                                              "encoded by CTG"
                                                                                                "encoded by GAC"
                                            "encoded
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0;

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produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
immunotherapeutic applications, including treatment of immune-mediated infections,
                                                                                                                                                                                                             genes can be expressed in a prokaryotic or eukaryotic host cells
                                                                                                                                                                                                                             transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface genes from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a more contained.
                                                                                                                                                                                                                                                                                                                                                                                                                                      were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by
                                                                                                                                                                                                                                                                                                                                                           in eukaryotic cells and selection of cells expressing adhesion to an antibody-coated substrate. The method the isolation and cloning of any protein which can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        efficient clohing and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This is the amino acid sequence of human deduced from cDNA clone p90 (see X07372) immunoselection cloning method from a cDN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 10; Column 53-54; 75pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    from a single patient undergoing extracorporeal induction therapy. Clones plas (see V63456) an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Plones plas (see v63456) and p98 (see x07373) A novel method for cloning cDNAs from mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cDNA library expressed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          I (FCRI),
                                                                                                                                                                                                                                                                                                                                                                                                          is useful
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W97834 standard; Protein; 374 Location/Qualifiers cell surface antigen; lymphocyte; human; "encoded by "encoded by TGG' 0; Ą Score 61; DB 19; Pred. No. 0.0016; Mismatches CTC" 0, Length 374; Indels 0, Gaps

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                                                            Query Match
Best Local Similarity
Matches 12; Conser
                                                                                                                                                                                                        expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V6344) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                           This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in CCC cells. The cDNA library was constructed from polyA RNA of cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-NOV-1998
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N-PSDB; X07373.
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                                                                                                                                                                                                                                                                                                                                                                                             from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones pl35 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 10; Column 53-54; 75pp; English.
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                                                                                                                                                                                                 immunotherapeutic applications,
                            1 EDGNVLKRSPEL 12
edgnvlkrspel 276
                                                                                                                                                                  including
                                                             Conservative
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88US-0160416.
89US-0379076.
90US-0553759.
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256 .
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176
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                                                       100.0%; Score 61; DB 19; 100.0%; Pred. No. 0.0016; rive 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Y96226
ID Y96226 standard; Protein;
                                                                                                                                         RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            eosinophil; HIV; IgG; immunosuppressive; antirheumatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y96226;
                                               B43683 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prevent autoimmune diseases, rheumatoid arthritis or MM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is the human high affinity FcgammaRI is also known as CD64. Fc receptors pl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection;
                                                                                                                                                                                                                                                                                   265
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                               Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fc receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 374;
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08-FEB-2001

(first entry)

Y96183 standard; Protein; 344.AA.

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C77607
B43398
                                                                                                                                                               Sequence
                                                                                                                                                                                                                antagonists may be also B44240 represent sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B43398 to B44239. The proteins can have activities based on and cells the genes are expressed in. Example of activities cytostatic; proliferative; vulnerary; immunomodulator; antic
                                                                                                                                                                                                                                                        inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, allergic reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cardiovascular disorders, neurological disease and bacterial or cancers, cardiovascular disorders, neurological disease and bacterial or cardiovascular disorders.
                                                                                                                                                                                                                                                                                                                                                    polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel isolated nucleic acids comprising sequences useful for treating or diagnosing e.g. cancer-
 290
                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-587533/55
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; cancer associated gene; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human cancer associated protein sequence SEQ ID NO:1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           allergic reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vasotropic; antipsoriatic; antiangiogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .mmune disorder;
                                 1 EDGNVLKRSPEL 12
edgnvlkrspel 301
                                                                       l Similarity
12; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      to C78448 encode the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 1739-1740;
                                                                                                                                                              399 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ruben SM;
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000WO-US05882.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thrombolytic; cardiovascular disorder; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antirheumatic; antiarthritic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiasthmatic;
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                                                                                                                                                                                                                sequences
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                                                                                                                                                                                                                            The peptides, nucleotides, antibodies, agonists and a also be used in drug screens. C78449 to C78457 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        aematopoietic cell disorder; autoimmune disorder; graft versus host disease; organ rejection;
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                                                                   Score 61; DB
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25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FcRI is a high affinity receptor for the Fc portion of IgG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FcRI and a receptor ligand will be helpful to increase
                               Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therap immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of from a cDNA library expressed in COS cells using a novel method of
                                                                                                Human macrophage-specific FCRI.
                                                                                                                                                                     Y96134;
                                                                                                                                                                                             Y96134 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment of immune-mediate animals, including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (panning on) an antibody-coated substrate such as a culture dish CSA nucleic acids isolated by the method of the invention, and the coated to the invention of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention designed to isolate CSA nucleic acids. The method based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
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                   amyloidosis;
                                                                                                                                  19-DEC-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                        Local Similarity 91.
les 11; Conservative
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| 65 edgmvlkrspel 276
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               parasitic disease;
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Stamenkovic I,
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N-PSDB; A50592.

Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic diseases.

Example 10; Column 53-55; 75pp; English.

The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids: The method is CD based upon transient expression of a CSA in eukaryotic cells and CC physical selection of cells expressing the antigen by adhesion to CC (panning on) an antibody-coated substrate such as a culture dish. CC proteins they encode, are useful for immunodiagnostic and CC immunotherapeutic applications, including the diagnostic and CC treatment of immune-mediated infections, diseases, and disorders in CC animals, including humans. These disorders include asthma, including humans. These disorders include asthma, collar complex disease, anyloidosis, parasitic diseases or multiple CC informally located on the cell surfaces of macrophages. The CC surfaces other than macrophages, is useful in therapy. A fusion CC protein of FCRI and a receptor ligand will be helpful to increase XXX.

Sequence 374 AA;

ν Query Match 86.9 Best Local Similarity 91.7 Matches 11; Conservative 86.9%; 91.7%; Score 53; DB 21; Length 374; Pred. No. 0.051; Indels 0; Mismatches 1; Indels 0; Gaps 0;

В 1 EDGNYLKRSPEL 12 ||| ||||||| |265 edgmvlkrspel 276

Search completed: June Job time: 205 sec 4, 2001, 12:13:32

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Result
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Query Match

82.0%;

Score 50;

DB 2;

Length 91;

US-08-332-562A-94 US-08-332-562A-94 US-08-332-562A-94 US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-32-562A-94  US-08-332-562A-94   ALIGNMENTS	28 33 54.1 480 4 US-09-170-670-5 29 33 54.1 480 4 US-09-193-068-5 30 33 54.1 480 4 US-09-183-412-5 31 33 54.1 480 5 PCT-US96-10521-9 32 33 54.1 503 3 US-08-258-287B-2 33 34.1 503 3 US-08-368-704C-2 35 33 54.1 503 3 US-08-368-704C-2 35 33 54.1 503 3 US-08-368-704C-2 36 33 54.1 503 3 US-08-368-704C-2 37 38 54.1 503 5 PCT-US93-05701-19 38 33 54.1 503 5 PCT-US93-05705-2 38 33 54.1 514 1 US-08-459-610-4 40 33 54.1 514 1 US-08-459-610-4 41 33 54.1 514 2 US-08-637-399-4 42 33 54.1 514 2 US-08-637-399-4 43 33 54.1 514 3 US-08-683-838A-4 44 33 54.1 514 3 US-08-683-838A-4 45 33 54.1 520 5 PCT-US94-01553A-34	
/08332562A  /08332562A  n F.C. k P. ret L. dette M. isa D. D. ph. ridner , N.W., Suite 50  isk atible boS/MS-DOS boS/MS-DOS boS/MS-DOS lease #1.0, Vers : : : : : : : : : : : : : : : : : : :	ALIGNMENTS	80 4 US-09-170-670-5 80 4 US-09-193-068-5 80 4 US-09-183-412-5 80 9 US-09-183-412-5 80 105-08-258-2878-2 80 3 US-08-258-2878-3 80 3 US-08-258-2878-3 80 3 US-08-368-704C-3 80 459-670-1 80 459-670-1 80 459-610-4 80 458-610-4 80 458-610-4 80 458-610-4 80 458-610-9088-4 80 458-700-9088-4 80 458-700-36 80 458-700-36 80 458-700-36 80 458-700-36
		Sequence 5, Ap Sequence 5, Ap Sequence 2, Ap Sequence 2, Ap Sequence 35, A Sequence 35, A Sequence 19, A Sequence 19, A Sequence 4, Ap Sequence 36, A Sequence 31, Ap

83.3%;

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RESULT 3
US-08-718-270A-11
                                             Sequence 11,
Patent No. 5
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Best Local
                              GENERAL INFORMATION:
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    APPLICANT:
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VEORMATION FOR SEQ ID NO:
                                                                                                                                200 EDGSVLKR 207
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TOPOLOGY: lir
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REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                         1 EDGNVLKR 8
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                                                       Application US/08718270A
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Hlavka, Joseph J.
Pincus, Matthew R.
                                                                                                                                                                                                                                                                                                                                                                          213 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Greenlee and Winner, P.C. 5370 Manhattan Circle, Suite
                                                                                                                                                                                                                                                                        Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hlavka, Joseph J.
Pincus, Matthew R.
No. 58406831e, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Abajian, Henry B. Kende, Andrew S.
                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                       protein
NO
                                                                                                                                                                                                                                                                                                                                             single
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N: 530
                                                                                                                                                                                                  59'.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptides Inhibiting the Oncogenic Action
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                                                                                                                                                                                                    Pred'. No.
                                                                                                                                                                                                                 Score
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                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suite 201
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                                                                                                                                                                                                              Length 213;
                                                                                                                                                                                  Indels
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Query Match
Best Local Similarity
"~+~hes 7; Conserv
                                                                                                                                                                                           Patent No.
                                                                                                                                                                                             Sequence 8, Application US/08737825 Patent No. 5871922
                                                                                                                                                                            GENERAL
                                                                                                                            APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (303) 499-808
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                   APPLICANT:
                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                      200 EDGSVLKR 207
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TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: No. 59104781e, APPLICANT: Abajian, Henry
                                                                                                                                                                                                                                                                                                       1 EDGNVLKR 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/004,091 FILING DATE: 21-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 08/531,525
FILING DATE: 21-SEP-1995
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                                                BYCROFT, BARRIE WALSHAM
WILLIAMS, PAUL
                                                                             COX; ANTHONY RICHARD JOHN HOLDEN, MATTHEW THOMAS GEOFFREY PORTER, LAUREN ELIZABETH
                                                                                                                              SEBAIHIA, MOHAMMED
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (303) 499-8080
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein
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GORDON SIDNEY ANDERSON BIRNIE
GENES INVOLVED IN THE BIOSYNTHETIC PATHWAY OF CARBAPENEM
12
                                                                                                                                           GEORGE PEACOCK COPELAND
SIMON JAMES
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Pred. No.
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13;
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ORRESPONDENCE ADDRESS:

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; ORIGINAL SOURCE:
; ORGANISM: Erw
US-08-737-825-8
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Best Local Similarity 70.v
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NFORMATION FOR SEQ ID NO:
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/394,189B
FILING DATE: 24-FEB-2005
                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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LENGTH: 288 amino acids
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 205-8000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               137 DGKTLKRVPE 146
                                                                                                                                                                                                                                         UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                     ITLE OF INVENTION:
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                                                                                                                                                          CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: SVENSSON, LEONARD R. REGISTRATION NUMBER: 30,330
                                                                                COMPUTER:
                                                                                                                                          COUNTRY:
                                                                                                                                                                                           STREET:
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                                                                                                                                                                                                          ADDRESSEE:
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Falls Church
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                                                                                                                                                                                             176 Federal Street
                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                      Shaham, Sha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (703) 205-8050
                                                                                                                                                                                                                                                                                                                   Yuan, Junying
                                                                                                                                                                                                                                                                                                                                     Horvitz, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Erwinia carótovora
                                                                             IBM Compatible
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                                                                                                                                                                                                          Clark & Elbing LLP
                                                                                             Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JUN-1997
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27
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70.0%;
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BETA CONVERTASE GENE TO A C. ELEGA
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Pred. No. 19;
0; Mismatches
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Best Local Similarity
6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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                                                                                                              FILING DATE: 24-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Bugaisky, Lawrence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 479 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/8
FILING DATE: 12-JUN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                             SOFTWARE: PatentIn Rel
                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                               PRIOR APPLICATION DATA:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-428-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                              ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   447 QGANILKQMPEL 458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ITLE OF INVENTION:
                           TELEFAX:
                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                              TELEPHONE:
                                                                           NAME: Bugaisky, Lawrence B. REGISTRATION NUMBER: 35,086 REFERENCE/DOCKET NUMBER: 06
                                                                                                                                              APPLICATION NUMBER: US 08/080,850 FILING DATE: 24-JUN-1993
                                                                                                                                                                                             FILING DATE: 10 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Bieker-Brady, Kristina REGISTRATION NUMBER: 39,109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    617-428-7045
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                                                                                                                                                                                                                                                                                                                                                              USA
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                                              (202) 371-2600
                                                                                                                                                                                                                10-JUN-1994
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ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Programmed Cell Death Genes and Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 57.4%;
50.0%;
                                                                                                                                                                                                                                                              Release #1.0, Version #1.25
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Pred. No.
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LENGTH: 497 amino acids

amino acid

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Best Local Similarity
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US-08-258-287B-37
                                                           Matches
                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                   NFORMATION FOR SEQ ID NO: 37:
                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 497 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tent No.
                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 35,086
REFERENCE/DOCKET NUMBER: 0609.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/080,850
FILING DATE: 24-UN-1993
ATTORNEY/AGENT INFORMATION:
465 QGANILKQMPEL 476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Yuan, Junying
APPLICANT: Miura, Masayuki
TITLE OF INVENTION: Programmed Cell Death Genes and Proteins
NUMBER OF SEQUENCES: 95
                                                                                                                                                                                                                                                                                                                                                                                                                     RIOR APPLICATION DATA:
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                                                                                                                                              TOPOLOGY:
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TELEX: 248636 SSK
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 0
FILING DATE: 10-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                            1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/368/704C FILING DATE: 4-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 EDGNVLKRSPEL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20005
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                                                                                                                                                                                                                                                                                                    Bugaisky, Lawrence B
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                                                        Conservative
                                                                                                                                             both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                               (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                4-JAN-1995
                                                                                                                                                                                                                             371-2540
                                                                       57.4%;
50.0%;
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                                                                    Score 35;
Pred. No.
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                                                     Mismatches
                                                                                                                                                                                                                                                                         .3920002
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                                                                              ДВ 3;
                                                                                Length 497;
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                                                     Indels
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RESULT

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US-08-368-704C-36
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                                                                                                                                                                                                                                                                                                                Sequence 36,
                                                                                                                                                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                           APPLICANT: Yuan, Junying APPLICANT: Miura, Masayuki TITLE OF INVENTION: Program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                            COUNTRY: USY
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PRIOR APPLICATION NUMBER: US 08/080,850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 24-JUN-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: PatentIn Rele
CURRENT APPLICATION DATA:
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TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                   471 QGSNILKOMPEL 482
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                            STREET: 1100 New CITY: Washington STATE: D.C.
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APPLICANT: Miura, Masayuki
NITLE OF INVENTION: Programmed Cell Death Genes and Proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity les 6; Conserv
                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                    1 EDGNVLKRSPEL 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 0609.3920001
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o. 6083735
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
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                                                                                                                                                         E: Sterne, Kessler, Goldstein & 1100 New York Avenue, Suite 600
                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     503 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Sterne, Kessler, Goldstein & 1100 New York Avenue, Suite 600
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; Pred. No. 56;
3; Mismatches
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US-08-394-189B-5
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Best Local Similarity
Matches 6; Conserv
                APPLICATION: 24-FEB-2001

CLASSIFICATION: 435

PRIOR APPLICATION NUMBER: 08/282,211

APPLICATION NUMBER: 07/984,182

APPLICATION NUMBER: 07/984,182

TTING DATE: 20-NOV-1992

TTING DATE: 20-NOV-1992

TTING DATE: 20-NOV-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      equence 5, Application US/08394189B atent No. 5962301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (202) 371-254
TELEX: 248636 SSK
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
LENGTH: 503 amino acid
                                                                                                                                                                                                                                               ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: US 08/258,287
                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
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471 QGSNILKQMPEL 482
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ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: 0609.3920002
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amino acid
GY: both
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                                                                                                                                                                                                                                                                                                                                                     176 Federal Street
                                                                                                                                                                                                                                                                                                      USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Horvitz, Robert
                                                                                                                                                                                                                               IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                         Clark & Elbing LLP
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лынк: 07/897,788
12-JUN-1992
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BETA CONVERTASE GENE TO A C. ELEGAN
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Pred. No. 56;
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Query Match
Best Local Similarity
Grands 6; Conserv
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Best Local Similarity 50.0
""" hes 6; Conservative
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SEQUENCE CHARACTERISTICS:
LENGTH: 505 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                          CLASSIFICATION:
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                OPENALE:
SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/05701
APPLICATION NUMBER: 19930614
                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 617-428-0200
                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
473 QGSNILKQMPEL 484
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                                                                                                                                                 NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 39,109
REFERENCE/DOCKET NUMBER: 01997/211001
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OPERATING SYSTEM:
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                          1 EDGNVLKRSPEL 12
                                                                                                                                                                                                              TYPE: amino acid
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                                                                                                                                                                                                                                            ENGTH:
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                                                                                                                                                                                                                                          505 amino acids
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NENTION: Cloning and Characterization of Cell Death Genes
                                                          Conservative
                                                                                                                                                 at every Xaa
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50.0%;
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                                                                       Score 35; DB Pred. No. 56;
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Pred. No. 56;
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                                                                                     Length 505;
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RESULT 12
PCT-US93-05705-5
REQUENCE 5, Application PC/TUS9305705
GENERAL INFORMATION:

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US-08-588-985-
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Best Local Similarity
Matches 6; Conserv
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Patent No. 5777094
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/I
FILING DATE: 19930714
INFORMATION FOR SEQ ID NO: 5.
SEQUENCE CHARACTERISTICS:
LENGTH: 505 amino acids
TYPE: amino acids
TYPE: amino acid
                   NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                             REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
FELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Warren M. Cheek, Jr.
NAME: Warren M. Cheek, Jr.
33,367
                                                                                                                                                                                                                                                    SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/588,985
FILING DATE: January 19, 1996
                                                                                                                                                                                                                                                                                                                                                                                                      STREET: busington CITY: Washington CTATE: D.C.
                                                                                                                                                                                                   CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25'inch, 500 kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
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TITLE OF INVENTION: Inhibitors of Ced-3 and Related Proteins
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                COMPUTER: IBM COMPONENTING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         473 QGSNILKQMPEL 484
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805 Fifteenth Street, N.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Michiyuki MATSUDA et al.
VENTION: CDNA OF DOCK180 GENE AND DOCK180 PROTEIN
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DN: 536
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Pred. No. 56;
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Best Local Similarity 66.7
Matches 8; Conservative
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ANTI-SENSE:
FRAGMENT TYPE:
ORIGINAL SOURCE:
US-08-971-988-2
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; ORIGINAL SOURCE:
US-08-588-985-2
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                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: WAITEN M. Cheek, Jr.
REGISTRATION UNUBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-8850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                          HYPOTHETICAL:
                                                                        MOLECULE TYPE:
                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 1865 amino acids
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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TITLE OF INVENTION:
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HYPOTHETICAL:
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                                                                                                   TYPE: amino acid
STRANDEDNESS: si
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                                                                                          TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
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TOPOLOGY: 11
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805 Fifteenth Street, N.W., #700
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ENTION: CDNA OF DOCK180 GENE AND DOCK180 PROTEIN
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Pred. No. 2.5e+02
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57.4%; Score 35; DB 1; 66.7%; Pred. No. 2.5e+02; tive 1; Mismatches 1

Length 1865;

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Gaps

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Search completed: June 4, 2001, 12:14:32 Job time: 225 sec
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                                                                                                                                                                                                                                              ; MOLECULE TYPE: protein
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
ORGANISM: Plasmodium falciparum
US-08-568-459A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 15
US-08-568-459A-12
                                                                                                                                       Query Match
Best Local Similarity 70.0
Thes 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: ISTACLSEN, Ned
REGISTRATION NUMBER: 29,655
REFERENCE/DOCKET NUMBER: NIHI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 235-850
TELEFAX: (619) 235-857
TELEFAX: (619) 235-0176
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 2710 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         equence 12, Application US/08568459A stent No. 5849306
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                                                                                     1680 DGNVLSVSPK 1689
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 3
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COUNTRY: US
ZIP: 92660
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FILING DATE: 07-DEC-1995
TRESTREAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Knobbe Martens Olson & Bear
STREET: 620 Newport Center Drive 16th Floor
CITY: Newport Beach
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: si
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Chitnis, Chetan
Miller, Louis H.
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Maximum Match 100%
Listing first 45 s
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Perfect score:
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2: pir2:*
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Fc gamma (IgG)
Fc gamma (IgG)
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replication protei
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succinate--CoA lig
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57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4	57.4
546	454	424	400	384	376	370	299	299	202	197	166	159	157	157	156
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C72453	A46498	JQ0661	T34363	T00625	S73941 .	C40585	D23696	S25782	T36138	S19102	T48040	B84096	T14381	S06074	T49264
hypothetical prote	glucocorticoid-sen	impB protein - Sal	hypothetical prote	, branched-chain ami	oligopeptide trans	recF protein - Sta	lacX protein - Lac	hypothetical prote	hypothetical prote	hypothetical prote		mutator MutT prote	heat-shock protein	heat shock protein	heat shock protein

## ALIGNMENTS

C; Species: Homo sapiens (man) C; Date: 03-Apr-1992 #sequence

#sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000; \$03019

FC gamma (IgG) receptor I (high affinity)
N;Alternate\_names: CD64

form b -

Science 243, 378-381,

Accession: A41357

R;Allen,

J.M.;

4135,, ; Seed, B. ; R1, 1989

C; Accession: A41357;

A; Title: Isolation and expression of functional high-affinity A; Reference number: A41357; MUID:89100284

Fc receptor complementa

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Fc gamma (19G) receptor I-A (high affinity) precursor - human
N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000
C;Accession: A39878; 170304; B41357; S03018; 157525
                                                                                                                                                                                                                                                                                                                          RESULT
A39878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 12; Conserva
                                               R;van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M. J. Biol. Chem. 266, 13449-13455, 1991
A;TItle: Gene organization of the human high affinity receptor for A;Reference number: A39878; MUID:91302383
A;Accession: A39878
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           F;117-170/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: EMBL:X14355, NID:g31333; PIDN:CAA32536.1; PID:g31334
A;Note: the authors translated the codon ACT for residue 25 as Ala
C;Superfamily: FC gamma receptor I; immunoglobulin homology
C;Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Molecule type: mRNA
A;Residues: 1-344 <ALL2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (A;Reference number: S03018; MUID:89098339
A;Accession: S03019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Allen, J.M.; Seed, B.
Nucleic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: mRNA
A;Residues: 1-344 <ALL1>
A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334
;Molecule type: DNA
;Residues: 1-374 <VAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                     265 EDGNVLKRSPEL 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 EDGNVLKRSPEL 12
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FC gamma (IgG) receptor high affinity - mouse
N; Alternate names: high affinity IgG receptor
C; Species: Mus musculus (house mouse)
C; Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change
                                                                                                           A; Reference number: A; Accession: A46480
                                                                                                                               A; Title: Structure and mapping of the gene encoding A; Reference number: A46480; MUID:92166399
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                                                                  Molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Reference number: 157525; MUID:93204964
A;Accession: 157525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Map position: 1q21-1q2; Superfamily: Fc gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: GB:S57204; NID:g298692;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;59,78,152,159,163,195,240/Binding site: carbohydrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: GB:M63830; R;Porges, A.J.; Redecha, P.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Keywords: alternative splicing; glycoprotein; 1-15/Domain: signal sequence #status predictec 16-292/Domain: extracellular #status predictec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                         Immunol. 148, 1570-1575,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Molecule type: DNA
Residuès: 1-8 <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Status:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             117-170/Domain: immunoglobulin homology <IMM2>
293-313/Domain: transmembrane #status predicted <TMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perez, C.; Wietzerbin, J.; Benech,
pl. Cell. Biol. 13, 2182-2192, 1993
Title: Two cis-DNA elements involve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Allen, J.M.; Seed, B.
cleic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Title: Isolation and expression of functional high-affinity Fc receptor complementary;Reference number: A41357; MUID:89100284;Accession: B41357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . Clin. Invest. 90, 210;Title: Novel Fc gamma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Residues:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        265
                                                                                                                                                                                                      N.; Kozak, C.A.; McKenzie, I.F.;
                                                         preliminary
e type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EDGNVLKRSPEL 12
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12; Conservative
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1-24,'8',26-337,'T',339-374 <AL1>
1-26,'8',26-337,'T',339-374 <AL1>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 61;
Pred. No.
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predicted <SIG>
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thes 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the human high affinity Fc receptor (Fc
                                                                                                                                                     mouse high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 374;
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                                                                                                                                               EC
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                                                                                                                                               RI
                                                                                                                                               and
                                                                                                                                                                        C; Accession: S77922
R; Dornelas, M.C.; Lecharny,
submitted to the EMBL Data
submitted to number: S77922
                                                                             A; Molecule type: mRNA
A; Residues: 1-407 < DOR>
                                                                                                                                                                                                                                                                                    shaggy-like protein kinase iota (EC 2.7.1.-)
C:Species: Arabidopsis thaliana (mouse-ear cr
C:)Date: 23-Jul-1997 #sequence_revision 01-Auc
        C; Keywords: ATP;
                   A;Cross-references: EMBL:X99696; NID:g1480077; PIDN:CAA68027.1; PID:g1480078
C;Superfamily: kinase-related transforming protein; protein kinase homology
                                                                                                                                                                                                                                                                                                                                                                               RESULT
S77922
                                                                                                                                       A;Status: preliminary
                                                                                                                                                                A; Accession: S77922
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F; 37-53/Domain:
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A; Map position: 12R
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-112 < DUZ>
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S55946
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phosphotransferase; protein
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4, 371-378,
                                                                                                                                                                                                                                  Lecharny,
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Kreis, M.

01-Aug-1997 #text\_change 20-Jun-2000

cress

Arabidopsis thaliana

August 1996

protein; in kinase

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probable membrane protein YLR390w - yeast (Saccharomyces N;Alternate names: hypothetical protein L8084.10 C;Species: Saccharomyces cerevisiae C;Date: 23-Aug-1995 #sequence_revision 19-Oct-1995 #text_C;Accession: S55946
                                                                                                                                                                                                 A;Cross-references: EMBL:U19729; NID:9625097;
A;Experimental source: strain S288C (AB972)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: GB:M31314; NID:g200752; PIDN:AAC;Superfamily: Fc gamma receptor I; immunoglobulin C;Keywords: immunoglobulin receptor; transmembrane F;127-179/Domain: immunoglobulin homology <IMM>
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R; Sears,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Title: Molecular cloning and expression A;Reference number: A43511; MUID:90111035
     Local Similarity 66.
nes 8; Conservative
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                                                                                                           transmembrane protein
                                                                                        transmembrane #status predicted
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                                                                                                                                                                                                                                                                                           sequence: S55944
                                                                                                                                               SGD:S0004382; MIPS:YLR390w
                      68.9%;
66.7%;
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1990
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B.; McKenzie,
                      Score 42;
Pred. No.
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Pred. No.
   Mismatches
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lobulin homology
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   2;
                                  Length 112;
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Gaps
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Best Local Similarity
Thes 8; Conserv
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Science 260, 695-698,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fc gamma (IgG) receptor high affinity – mouse
C;Species: Mus musculus (house mouse)
C;Sate: 02-Jul-1996 #sequence_revision 02-Jul-1996 #text_change 21-Jan-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               F; 68-329/Domain: F; 76-84/Region: F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        conserved hypothetical protein VC0302 [imported] - Vibrio cholerae (group Ol
                                                                                                                                                                                                                                                                                                                                                                                  l, R.R.; Mekalanos, J.J.;
Vature 406, 477-483, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Molecule type: mRNA; Residues: 1-336 <RE
                                                                                                                                                                                                                                                                          ; Molecule type: DNA
; Residues: 1-396 <H
                                                                                                                                                                                                                                                                                                                             ;Title: DNA Sequence of both chromosomes of the cholera pathogen Vibrio cholerae.;Reference number: A82035; MUID:20406833;Accession: H82339
                                                                         Best Loc
Matches
                                                                                                                                                                                                     ;Gene: VC0302
                                                                                                                                                                                                                                   ;Cross-references: GB:AE004119; GB:AE003852; NID:g9654718; PIDN:AAF93476.1; GSPDB:GN00;
Experimental source: serogroup Ol; strain N16961; biotype El Tor
                                                                                                                                                                                                                                                                                                                                                                                                    Heidelberg, J.F.; Eisen, J.A.; Nelson, W.C.; Clayton, hardson, D.; Ermolaeva, M.D.; Vamathevan, J.; Bass, S., R.R.; Mekalanos, J.J.; Venter, J.C.; Fraser, C.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reference number: I48471; MUID:93242399
                                                                                                                                                                                                                        Genetics
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title: Linkage on chromosome 3 of autoimmune diabetes and defective Fc receptor for
                                                                                                           Query Match
                                                                                                                                                                                  Map position: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Species: Vibrio cholerae
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   383 QDGNALKAEPE 393
                                                                       Local Similarity 63.6 hes 7; Conservative
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                                     EDGNVLKRSPE 11
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protein kinase ATP-binding motif
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66.7%;
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72.7%;
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Pred. No.
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Pred. No.
                                                                                        Score 40; DB
Pred. No. 9.7;
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4.2;
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5.2
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Bass, S.; Qin, H.; Dragoi, I.; Sellers,
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C;Species
replication protein - Bacillus subtilis plasmid pIM13 C;Species: Bacillus subtilis C;Date: 05-Oct-1988 #sequence_revision 05-Oct-1988 #t
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A25233
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Best Local Similarity
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A;Gene: APE1307
C;Superfamily: Synechocystis long-chain-fatty-acid--CoAC;Superfamily: Synechocystis long-chain-fatty-acid--CoAC;Keywords: acid-thiol ligase; ccenzyme ACC;Keywords: acetate--CoA ligase homology <ACL>
                                                                                                                                                                                                                                                            A;Cross-references: DDBJ:AP000061; NID:g5104821; PIDN:BAA80298.1; PID:g5104984
A;Experimental source: strain Kl
                                                                                                                                                                                                                                                                                                 A;Molecule type: DNA
A;Residues: 1-651 <KAW>
                                                                                                                                                                                                                                                                                                                                                        A; Reference number: A; Accession: D72605
                                                                                                                                                                                                                                                                                                                                                                       A;Title: Complete genome sequence of an aerobic hyper-thermophilic Crenarchaeon, Aero A;Reference number: A72450; MUID:99310339
                                                                                                                                                                                                                                                                                                                                                                                                            R;Kawarabayasi, Y.; Hino, Y.; awa, H.; Takamiya, M.; Masuda, DNA Res. 6, 83-101, 1999
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C; Accession:
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Best Local S
Matches 7
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432
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EDGEILMRSPAM
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                                                                       Conservative
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M.; Masuda, S.; Funahashi,
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443
                                 12
                                                                                         58.3%;
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Pred.
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                                                                                         NO;
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                                                                                                          Length 651
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T.; Kudoh, Y
                                                                                                                                                                                                      acetate--CoA ligase
                                                                       0;
                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                .; Jin-no, K.;
Y.; Yamazaki,
                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Aeropy
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Genetics 148, 1021-1029, 1998
A; Title: Theme and variation among silencing proteins in A; Reference number: %16686; MUID:98198828
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; Rine, U.
R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, S.U.; R;Astrom, 
A;Description: involved in both silencing C;Keywords: transcription regulation
                                                                                                                                                                                                                                                                                                                                                                                                           A; Cross-references: EMBL: AF035007; NID: g3002787; A; Note: the source is designated as Kluyveromyce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N; Alternate names: silent information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mating type silencing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Accession: T09481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;Species: Kluyveromyces marxianus
                                                                                                                                                                                                                                                                                                                                                              Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Residues: 1-1314 <AST>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                   Function:
                                                                                                                                                                                                                                                           SIR4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-Sep-1999 #sequence_revision 20-Sep-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T09481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       regulator SIR4 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      regulator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       yeast (Kluyveromyces marxianus var. lactis)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .actis, Candida sphaerica
                                                                                and telomere length maintenance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PID: g3002789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            #text_change 11-May-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Saccharomyces cerevisiae and
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EDGNVSEKTPQ 717 EDGNVLKRSPE 11

revision 05-Oct-1988 #text\_change 10-Dec-1999

Conservative

62.3%; 54.5%;

Score 38; Pred. No.

84; 2

Length 1314,

0

Gaps

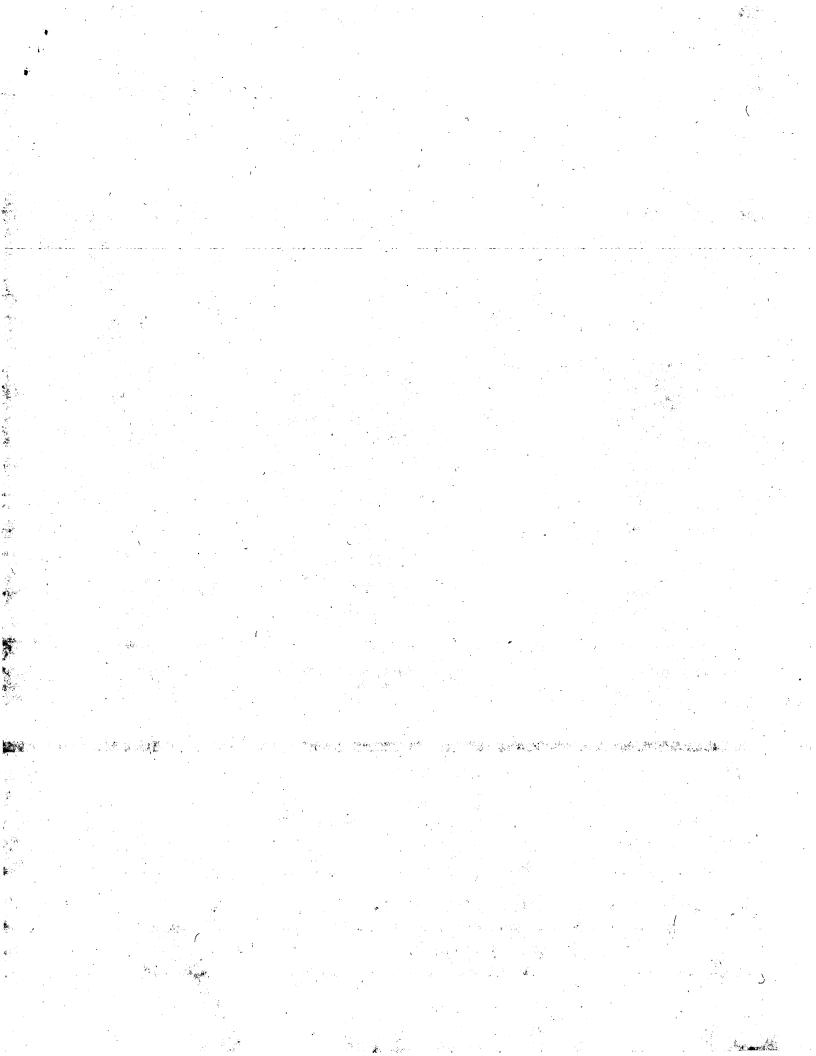
Mismatches

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C;Accession: A61069
R;Dyke, K.G.H.; Curnock, S.P.
FEMS Microbiol. Lett. 58, 209-216,
                                                                                     C;Species: Staphylococcus aureus
C;Date: 31-Dec-1993 #sequence_revision
             A: Accession:
                       A; Title: The nucleotide A; Reference number: A61(
                                                                                                                                                                                    B
                                                                                                                  replication
                                                                                                                                                                                                                                                                                                    C; Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Accession: A R; Monod, M.; D J. Bacteriol.
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                                                                                                                                                                                                                                                                                                                                                                                  N; Molecule type: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  replication
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: Sequence and properties of pIM13, A;Reference number: A91819; MUID:86250583 A;Accession: A25233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-146 <MON>
                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                         Experimental source:
                                                                                                                                                                                                                                                                                                                                                                    Residues: 1-158 <CAT>
                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                            Status: preliminary; not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Molecule type: DNA PASSidues: 1-158 <PRO>
                                                                                                                                                                                                                                                                                                                                            Comment: This sequence
                                                                                                                                                                                                                                                                                                                                                                                                                     Reference number: A46568; MUID:89036120
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Experimental source: plasmid pE5 Catchpole, I.; Thomas, C.; Davie, Gen. Microbiol. 134, 697-709, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cross-references: GB:M17990; NID:g153063; PIDN:AAA98227.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Title: Replication properties of pIM13, Reference number: A91847; MUID:88032835 Accession: A29827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Date: 18-Oct-1989 #sequence_revision 18-Oct-1989 #text_change 10-Dec-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Accession: A29827; C46568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Species: Staphyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                         resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteriol.
                                                                                                                                                                                    101
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                                                                                                                                                                                                                                     Local Similarity
les 8; Conserv
                                                                                                                                                                                                       1 EDGNVLKR-----SPEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                           The nucleotide sequence of Staphylococcus aureus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EEGNIIKRKTGVLMLNPEL 119
                                                                                                             protein REP - Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein REP - Staphylococcus aureus
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8; Conserv
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. 167, 13
                                                                                                                                                                                                                                                                                                     plasmid
                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monod,
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                       A61069
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138-147, 1986
                                                                                                                                                                                                                                                                                                replication protein
                                  sequence
                                                                                                                                                                                                                                                                                                                                                       plasmid
                                                                                                                                                                                                                                             61.5%;
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42.1%;
                                                                                                                                                                                                                                                                                                                                                                                        compared with conceptual translation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Davies,
                                                                                                                                                                                                         12
                                                                                                                                                                                                                                                                                                                                                     pT48
                                 of a
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1988;
1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                     the only long
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                                                                                                                                                                                                                                 Mismatches
                                  small
                                                                                   31-Dec-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C.S.;
                                                                                                                                                                                                                                               37.5;
No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a naturally occurring
                                                                                                                                                                                                                                                                                                 REP
                                cryptic
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                                                                                                             plasmid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plasmids
                                                                                                                                                                                                                                                                                                                                     open
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                                                                               #text_change 10-Dec-1999
                                                                                                                                                                                                                               ,0
                              plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2;
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                                                                                                                                                                                                                                                       Length 158;
                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                    reading
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 146;
                                                                                                           pox1000
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                                                                                                                                                                                                                                                                                                                                                                                                                                           plasmid
                              found
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PID: g153064
                                                                                                                                                                                                                                                                                                                                     frame
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid found
                                                                                                                                                                                                                            7;
                          in Staphylococcus aur
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7;
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                                                                                                                                                                                                                                                                                                                                    of a
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RESULT
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R;Keen,
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S19534
                                                                                                                  C; Superfamily: lipocalin;
                                                                                                                                                                                                                                       crustacyanin chain C1 - Euro
C;Species: Homarus gammarus
C;Date: 19-Mar-1997 #sequenc
                                                                                                                              A; Molecule type: protein A; Residues: 1-181 < KEE>
                                                                                                                                                                   A; Title: Complete sequence and model for the C(1) subunit of A; Reference number: S19534; MUID:92037648
                                                    Query Match
Best Local S
Matches 7
                                                                                                                                                            A; Accession:
                                                                                                                                                                                                   Eur. J. Biochem.
                                                                                                                                                                                                                              C; Accession: S19534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-162 < LAM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: A; Accession: B24497
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A;Molecule type: DNA
A;Residues: 1-158 <DYK>
C;Comment: This sequence represents
                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C; Superfamily: plasmid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₹;Lampson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   replication protein - Staphylococcus epidermidis plasmid pNE131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;Title: Nucleotide sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Species: Staphylococcus epidermidis Date: 05-Jun-1988 #sequence_revision
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and pSN2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lampson, B.C.; Parisi, J.T.
Bacteriol. 167, 888-892, 1986
73 DGNLLKRNGKL
                                                                                                                                                                                                                                                                                                                                     101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          101 EEGNIIKRKTGALMLNPEL 119
                       2 DGNVLKRSPEL 12
                                                                                                                                                                                                             J.N.; Caceres,
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                                                                                                                                                                                                                                                                                                                                                             EDGNVLKR----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plasmid
                                                     Similarity 7; Conserva
                                                                                                                                                            S19534
                                                                                                                                                                                                                                                                                                                                                                                       Similarity 42.3
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B24497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 42.
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      plasmid replication
                                                    Conservative
                                                                                                                                                                                                   202,
                                                                                                                                                                                                                                     #sequence_revision
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                                                                                                                                                                                                                                                  - European lobster
mmarus (European lo)
                                                                                                                                                                                                  31-40, 1991
                                                                                                                                                                                                           I.; Eliopoulos, E.E.;
                                                                                                                                                                                                                                                                                                                                                            ---SPEL 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            replication
                                                               60.7%;
                                                                                                             lipocalin homology
                                                                                                                                                                                                                                                                                                                                                                                                  61.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the constitutive
                                                                          Score
                                                                                                                                                                                                                                                                                                                                                                                    Score 37.5; I
Pred. No. 11;
4; Mismatches
                                                               Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein
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Pred. No. 1
                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the only long open
                                                                                                                                                                                                                                    lobster)
| 25-Apr-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                         37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-Jun-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          translation
                                                           DB
15;
                                                                                                                                                                                                      Zagalsky, P.F.; Findlay, J.B.C
                                                                                                                                                                                                                                                                                                                                                                                                              DВ
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Maximum Match 100%
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Gapop 10.0 , Gapext 0.5
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YBAZ_ECOLI
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vibrio harv
schizosacch
                  homo sapien
rattus norv
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P12314; P12315;
01-OCT-1989 (Rel. /12, Created)
01-OCT-1989 (Rel. 12, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I PRECURSOR (FC-GAMMA RI) (FCRI) (IGG FC RECEPTOR I) (CD64).

FCGRIA OR FCGRI OR FCGI OR IGFRI.
HOMO Sapiens (Human).
EUKaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Bukaryota; Metazoa; Chordata; Craniata; Hominidae; Homo.
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YF92_SYNY3	YGHR_ECOLI	SC4_OCTDO	VP11_RDVF	VP11_RDVA	HS11_WHEAT	VGLP_BEV	NAH1_YEAST	YKFO_YEAST	L100_ADEGX	KIRI_SALTY	KIRI_ECOLI
		P27012 octopus dof							P36856 avian adeno		P08204 escherichia

## ALIGNMENTS

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EMBL; X14356; CAA32537.1; EMBL; X14355; CAA32536.1; PIR; S03018; S03018. DTB: S03019; S03019	RO S B p t t ui	MEDLINE-89100284; pubMed=2911749; Allen J.M., Seed B.; "Isolation and expression of functional high-affinity Fc receptor complementary DNAs."; Science 243:378-381(1989)I- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS GAMMA. HIGH AFFINITY RECEPTOR.	SEQUENCE FROM N.A.  MEDLINE-89098339; PubMed-2974947;  Allen J.M., Seed B.;  "Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (FCRI).";  Nucleic Acids Res. 16:11824-11824(1988).  [2]  SEQUENCE FROM N.A.
SWISS-PROT entry is copyright. It is produced through a collaborat ween the Swiss Institute of Bioinformatics and the EMBL outstatic European Bioinformatics Institute. There are no restrictions on by non-profit institutions as long as its content is in in ified and this statement is not removed. Usage by and for commercities requires a license agreement (See http://www.isb-sib.ch/annousiend an email to license@isb-sib.ch).			MEDILINE-89100284; PubMed=2911749; Allen J.M., Seed B.; *Isolation and expression of functional high-affinity Fc recept complementary DNAS."; Science 243:378-381(1989)1- FUNCTION: BINDS TO THE FC REGION OF IMMUNOGLOBULINS GAMMA. AFFINITY RECEPTOR.

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RESULT
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P26151;
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01-MAY-1992
15-JUL-1999
                                           Osman N., Kozak C.A., McKenzie I.F., Hogarth P.M., "Structure and mapping of the gene encoding mouse gamma RI and chromosomal location of the human Fc J. Immunol. 148:1570-1575(1992).
                                                                               SEQUENCE FROM N.A. MEDLINE=92166399;
                                                                                                                                                                                                                                                                                                                                                                                         CONFLICT
SEQUENCE
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Mammalia; Eutheria;
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CARBOHYD
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CARBOHYD
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                                                                                                                    "Molecular cloning and
                                                                                                                                 MEDLINE=90111035; PubMed=2136886;
Sears D.W., Osman N., Tate B., Mc
                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                              Mus musculus
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Immunoglobulin
                                                                                                          Immunol.
        FUNCTION: BIRDS TO THE FC REGION OF IMMUNOGLOI AFFINITY RECEPTOR.
SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN TISSUE SPECIFICITY: MACROPHAGE-SPECICIC.
                                                                                                                                                                                                                                                                                                                      1 EDGNVLKRSPEL
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CONTAINS 3 IMMUNOGLOBULIN
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                                                                              PubMed=1531670;
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Rodentia;
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Last annotation
LOBULIN GAMMA FC
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IMMUNOGLOBULIN-LIKE C2-TYPE
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N-LINKED
                                                                                                                                                                         Craniata; Vertebrata; | Sciurognathi; Muridae;
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S -> T (IN CAA32536).
2C2AA8103ECF16E6 CRC64;
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
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Pred. No.
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                                                                                                                       Hogarth P.M.; high affinity
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01-OCT-2000
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01-OCT-2000
                          STRAIN=CV. COLUMBIA;
Piao H.L., Jang H.J.,
                                                                         Dornelas M.C., Schwebel-Dugue N., Thomas "Three new cDNAs related to SGG/GSK-3 (SI Kinase-3) from Arabidopsis thaliana.";
                                                                                                                                                                                          01-OCT-2000 (Rel. 40, 1
01-OCT-2000 (Rel. 40, 1
SHAGGY-RELATED PROTEIN
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Dornelas M.C., Schwei
                                                                                                                                            Eukaryota; Viridiplantae; Embryophyta; Trache Magnoliophyta; eudicotyledons; core eudicots; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                          ASK9 OR GSK1 OR 120000.
Arabidopsis thaliana (Mouse-ear cress).
Arabidopsis thaliana (Mouse-ear cress).
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SEQUENCE
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                                                                                                                                                                                                                                                                                                                             Similarity 83.
10; Conservative
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                                                                                                                                                                                                                                        STANDARD;
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                         Pih K.T., Lim J.H., Kang S.G.,
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EXTRACELLULAR (POTENTIAL)
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IG-LIKE C2-TYPE DOMAIN.
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                                                                                 as M., Lecharny A., Kreis (SHAGGY/glycogen synthase
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e; eurosids II;
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                                                                                            Kreis M.;
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Best Local :
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DOMAIN 70
NP_BIND 76
BINDING 99
ACT_SITE 195
SEQUENCE 407 AA
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Altafi H., Brooks S., Buehler E., Chao Q., Con
Altafi H., Brooks S., Buehler E., Chao Q., Con
Hansen N.F., Johnson-Hopson C., Khan S., Kim C
Hansen N.F., Southwick A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL;
                      SEQUENCE FROM N.A. MEDLINE=86250583; Puk Monod M., Denoya C.,
                                                                                                                                                                                                                             BACSU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=CV. COLUMBIA;
Liu S.X., Vaysberg M.
macrolide-lincosamide-streptogramin
                                                                        Bacillus/Staphylococcus
NCBI_TaxID=1423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pfam; .PF00069; pkinase; 1.
PROSITE; PS00107; PROTEIN_KINASE_ATP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or send an email to license@isb-sib.ch).
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                                                                                                   Bacteria; Firmicutes;
                                                                                                             Plasmid pIM13
                                                                                                                           Bacillus subtilis.
                                                                                                                                                   REPLICATION
                                                                                                                                                                01-JAN-1990
15-DEC-1998
                                                                                                                                                                                        01-JAN-1990
                                                                                                                                                                                                     P13968;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mendel; 13957; Arath;2319;13957
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European Bioinformatics Institute. There a
by non-profit institutions as long as i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE SER/THR FAMILY OF PROTEIN KINASES CDC2/CDKX SUBFAMILY; GSK-3 SUBSUBFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nitted (JUL-2000) to the EMBL/GenBank/DDBJ databases. FUNCTION: MAY MEDIATE EXTRACELLULAR SIGNALS TO REGULATE TRANSCRIPTION IN DIFFERENTIATING CELLS (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BY SIMILARITY)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P24941; 1AQ1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AF019927; AAB71545.1; AC068143; AAF82167.1;
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                                                                                                                                                                                                                                                                                                                              Similarity 72.
8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PS00108; PROTEIN_KINASE_ST; PS50011; PROTEIN_KINASE_DOM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AUTOPHOSPHORYLATED MAINLY ON THREONINE AND SERINE RESIDUES
            Denoya C., Dubnau D.; and properties of pIM13, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IPR002290; -.
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                                                                                                                                                   AND MAINTENANCE PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Serine/threonine-protein kinase;
                                                                                                                                                                                                                  STANDARD;
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                                      PubMed=3087948;
                                                                                                                                                                                                                                                                                                                                                                                           46024 MW;
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Last annotation updat
                                                                                                                                                                                       Created)
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                                                                                     group; Bacillus.
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د.
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BY S
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Pred. No. 1.6;
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1; Mismatches
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96BC4C53754A764C CRC64;
 B resistance plasmid
                                                                                                                                                  (PLASMID REPLICATION PROTEIN)
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Smith A.,
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or send a
PIR; AZYOL.,
DNA replication; PI
DNA TENDER 158 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Staphylococcus aureus, and Staphylococcus simulans Plasmid pE5, and Plasmid pPV142.
Bacteria; Firmicutes; Bacillus/Clostridium group;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; M13761; AAA98135.1; PIR; A25233; A25233.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bacillus subtilis.";
J. Bacteriol. 167:138-147(1986)
                                                                                                                                                                                                                                                                                                     6
                                                               EMBL; M17990; AAA98227.1; -.
                                                                                             or send an email to license@isb-sib.ch).
                                                                                                            entities requires a
                                                                                                                        use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                     the European Bioinformatics Institute.
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                                                                                                                                                                                                              Somkuti G.A., Solaiman D.K.Y., Steinberg D.H.; Submitted (OCT-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                           SPECIES-S
                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                               "Replication properties of pIM13, a naturally occurring plasmid found in Bacillus subtilis, and of its close relative pE5, a plasmid native
                                                                                                                                                                                                                                                                                                                                                           SPECIES=S.aureus;
MEDLINE=88032835;
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                Bacillus/Staphylococcus group;
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                                                   EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=1280,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       101 EEGNIIKRKTGVLMLNPEL 119
                                                                                                                                                                                                                                                                                    Staphylococcus aureus.";
Bacteriol. 169:5131-5139(1987).
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                                                 AF019140;
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8; Conservative
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                                                                                                                                                                                                                                           .simulans; STRAIN=13044; PLASMID=pPV142;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Rel.
                                    A29827
                                                                                                                                                                   Swiss Institute of Bioinformatics
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                                                   AAC33147.1;
                     Plasmid
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13,
37,
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       18079 MW;
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42.1%;
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Last annotation updat
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         60E337A2A024455C CRC64;
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Query Match
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61.5%;

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O1-AUG-1991 (Rel. 19, Created)
O1-AUG-1991 (Rel. 19, Last sequence up
O1-CCT-1994 (Rel. 30, Last annotation
CRUSTACYANIN C1 SUBUNIT
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    "Complete sequence carotenoprotein, cr
                                                   Findlay J.B.C.;
                                                                                                                                                               Nephropoidea; Nephropidae; Homarus NCBI_TaxID=6707;
                                                                                                                                                                                                           Homarus gammarus (European lobster) (Homarus vulgaris).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Astació
                                                                                                                                                                                                                                                                                                                                                                                                CRC1
                                                                                                  MEDLINE=92037648;
                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                          Homarus gammarus (European lobster)
                                                                                                                                                                                                                                                                                                                                                                                                                   HOMGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lampson B.C., Parisi J.T.;
"Nucleotide sequence of the constitutive macrolide-lincosamide-
"Nucleotide sequence of the constitutive macrolide-lincosamide-
streptogramin B resistance plasmid pNE131 from Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria; Firmicutes; & Bacillus/Staphylococcus
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01-JAN-1988 (Re
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MEDLINE=86304180; PubMed=3091582;
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UENCE 162 AA; 18447 MW;
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                                                                           J.N.,
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8; Conser
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    crustacyanin,
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06, Last sequence update)
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37, Last annotation update)
                                                                   PubMed=1935978;
s I., Eliopoulos
    and model for the rustacyanin, and mc
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01-APR-1988
01-APR-1988
01-OCT-2000
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                                                                 MEDLINE=97661202: PubMed-8905232:
OShima T., Aiba H., Baba T., Pujita K., Hayashi K., Honjo A.,
Ikemoto K., Inada T., Itoh T., Kajihara M., Kanai K., Kashimoto K.,
Ikemoto K., Inada T., Honto T., Kajihara M., Kanai K., Mashimoto K.,
Kimura S., Kitagawa M., Makino K., Masuda S., Miki T., Mizobuchi K.,
Mori H., Motomura K., Nakamura Y., Nashimoto H., Nishio Y., Saito N.,
Sampei G., Seki Y., Tagami H., Takemoto K., Wada C., Yamamoto Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.

MEDLINE-86104124; PubMed-3002435;
Buck D., Spencer M.E., Guest J.R.;

"Primary structure of the succiny'
coli.";
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                         Yano M., Horiu
"A 718-kb DNA
                                                                                                                                                                                                                                                                                       Mau B., Shao Y.; "The complete genome sequence of Escherichia coli K-12."; Science 277:1453-1474(1997).
                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-97426617; PubMed-9278503;
Blattner F.R., Plunkett G. III, Bloch C.A.,
Riley M., Collado-Vides J., Glasner J.D., R
Gregor J., Davis N.W., Kirkpatrick H.A., Go
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Pfam; PF00061; lipocalin; 1.
PRINTS; PR01273; INTERTCOLOR.
PROSITE; PS00213; LIPOCALIN; FALSE_NEG.
                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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STRAIN=K12 / MG1655;
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TYPES OF SUBUNITS: A1, A2, A3, C1,
-i- SUBCELLULAR LOCATION: EXTRACELLULAR;
-i- SIMILARITY: BELONGS TO THE LIPOCALIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COLORATION TO THE CARAPACE OF THE LOBSTER.
-I- SUBUNIT: OLIGOMER; CAN FORM DIMERS (BETA-CRUSTACYANIN);
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                                                  Horiuchi T.;
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51
181 AA;
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(Rel. 07, Last sequence update)
(Rel. 40, Last annotation update)
A SYNTHETASE BETA CHAIN (EC 6.2.1.5)
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  sequence of the Escherichia to the 12.7-28.0 min region
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IDS THE CAROTENOID
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Pred. No.
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coli K-12 genome on the linkage m
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Biochemistry 39:17-25(2000).
                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
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J. Biol. Chem. 269:10883-10890(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=99141407; PubMed=9917402; Fraser M.E., James M.N., Bridger W.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wolodko W.T., Fraser M.E., James M.N.G., Bridger Wolodko W.T., Fraser M.E., James M.N.G., Bridger W
                                                                                                                                                         or send
                                                                                                                                                                         entities
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "A detailed structural description of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           X-RAY CRYSTALLOGRAPHY (2:5 ANGSTROMS).
MEDLINE=94193797; PubMed=8144675;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Electrophoresis 18:1259-1313(1997).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=K12 / EMG2;
MEDLINE=97443975; PubMed=9298646;
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SWISS-2DPAGE; P07460; COLI.
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in the genome of Escherichia coli K-12.";
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                                                                                                                                                                                                                                                                      SUCCINATE. THE COMPLETE ACTIVE SITE IS PROBABLY LOCATED REGION OF ALPHA- BETA CONTACT.
SIMILARITY: TO OTHER BETA.
                                                                                                                                                                                                                                                                                                                                                         CITRIC ACID CYCLE, COUPLING THE HYDROLYSIS OF SUCCINYL-COA TO THE SYNTHESIS OF ATP & THUS REPRESENTS AN IMPORTANT SITE OF SUBSTRATE-LEVEL PHOSPHORYLATION. IT CAN ALSO FUNCTION IN THE OTHER DIRECTION FOR ANABOLIC PURPOSES, AND THIS MAY BE PARTICULARLY IMPORTANT FOR PROVIDING SUCCINYL-COA DURING ANAEROBIC GROWTH WHEN THE OXIDATIVE ROUTE FROM 2-OXOGLUTARATE IS SEVERELY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBUNIT: TETRAMER OF TWO ALPHA AND TWO BETA CHAINS.
MISCELLANEOUS: SUCCINYL-COA SYNTHETASE (SCS) OF E.COI
MISCELLANEOUS THREE STEPS THAT INVOLVE PHOSPHORYL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ENZYME REGULATION: EXHIBITS TWO INTERESTING PROPERTIES: "SUBSTRATE SYNERGISM", IN WHICH THE ENZYME IS MOST ACTIVE FOR THE CATALYSIS OF ITS PARTIAL REACTIONS ONLY WHEN ALL THE SUBSTRATE BINDING SITES ARE OCCUPIED, AND "CATALYTIC COOPERATIVITY" BETWEEN ALTERNATING ACTIVE SITES IN THE TETRAMER, WHEREBY THE INTERACTION OF SUBSTRATES (PARTICULARLY ATP) AT ONE SITE IS NEEDED TO PROMOTE
                                                                                                                                                                                                                   European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENZYME-BOUND SUCCINYL PHOSPHATE AS INTERMEDIATES.
MISCELLANEOUS: DURING AEROBIC METABOLISM IT FUNCTIONS IN THE
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                                                                                                           AE000176;
                                                                                                                        J01619; AAA23899.1;
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                                                                                                                                                                    and this statement is not removed requires a license agreement (See
                                                                                                                                                                                                     non-profit
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                                                                                                                                                         email to license@isb-sib.ch).
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                             0-JAN-00
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                                                                                           BAA35394.1;
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TO OTHER BETA SUBUNITS OF SUCCINYL-COA SYNTHETASES.
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01-JUL-1993
01-JUL-1993
                                                            Heat shock; SEQUENCE 1
                                                                                                                                                                                                                                                                                                                                Nishi R., Hashimoto H., Uchimiya H., Kato A.; Submitted (JUL-1992) to the EMBL/GenBank/DDBJ-I-SUBUNIT: FORMS OLIGOMERIC STRUCTURES.
-I-SUBCELLULAR LOCATION: CYTOPLASMIC.
-I-SIMILARITY: BELONGS TO THE SMALL HEAT SHOO
                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=4530;
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P31673;
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PROSITE; PS01217; SUCCINYL_COA_LIG_3; l.
Ligase; Tricarboxylic acid cycle; 3D-structure.
SEQUENCE 388 AA; 41392 MW; 09C429EC97A823CF CRC64;
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EcoGene; EG10981; s
                                                                                                                            EMBL; D12635; BAA02160.1; PIR; JS0710; JS0710.
                                                                                                                                                                                        modified and this statement
                                                                                                                                                                                                                                                                                                                                                                                                                                                Magnoliophyta; Liliopsida;
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Eukaryota; Viridiplantae;
                                                              Pfam, PF00011; HSP20; 1.
PROSITE; PS01031; HSP20; 1.
Heat shock; Multigene family.
SEQUENCE 154 AA; 17376 MW;
                                                                                                                                                                 or send
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                                                                                                                                                                                                      use
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                                                                                                               InterPro; IPR002068;
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8; Conserv
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I HEAT SHOCK PROTEIN
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; Poales; Poaceae; Ehrhartoideae; Oryzeae;
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RAS-RELATED
                                           Eukaryota; Viridiplantae; Embr.
Magnoliophyta; eudicotyledons;
Brassicales; Brassicaceae; Ara
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SEQUENCE
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Multigene family; Transcription regulation.
DOMAIN 20 83 KRAB RELATED.
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15-DEC-1998 (Rel
15-DEC-1998 (Rel
01-OCT-2000 (Rel
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Int. J. Cancer 72:965-971(1997).
-!- FUNCTION: COULD ACT AS A MODULATOR OF TRANSCRIPTION.
-!- SIMILARITY: BELONGS TO THE SSX FAMILY.
-!- SIMILARITY: CONTAINS 1 KRAB-RELATED DOMAIN.
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Gure A.O., Tureci O., Sahin
Knuth A., Pfreundschuh M., O
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               de Leeuw B., Balemans M., Geurts van Kessel A., "A novel Kruppel-associated box containing the SSX gene (SSX3) human X chromosome is not implicated in t(X;18)-positive synov:
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U90840; AAC05819.1;
TE; PS50806; KRAB_REI
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PubMed=8697803;
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Catarrhini; Hominidae
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                                                              Rosidae;
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GFAP_E
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SEQUENCE OF 197-227 FROM N.A.

SEQUENCE OF 197-227 FROM N.A.

Kirkpatrick B.W., Hart G.L., Moore S.S., Barendse W.;

Submitted (OCT-1993) to the EMBL/GenBank/DDBJ databases.

'- FUNCTION: GRAP, A CLASS-III INTERNEDIATE FILAMENT, IS A CELL-
SPECIFIC MARKER THAT, DURING THE DEVELOPMENT OF THE CENTRAL

NERVOUS SYSTEM, DISTINGUISHES ASTROCYTES FROM OTHER GLIAL CELLS.
      This SWISS-PROT
                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                          TISSUE=Subcommissural organ;
Bouchard P.;
                                                                                                                                                                                                                                      NCBI_TaxID=9913;
                                                                                                                                                                                                                                                     Bovidae; Bovinae;
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                                                                                                                                                                                                        SEQUENCE FROM N.A.
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PIR; JS0639; JS0639.
HSSP; P28746; 1A2K.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Isolation and analysis of
of Arabidopsis thaliana.";
Gene 108:259-264(1991).
-!- SIMILARITY: TO RAS PROT
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InterPro; IPR002380; -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=VARIOUS STRAINS; TISSUE=Leaf;
MEDLINE=92084144; PubMed=1748311;
Anai T., Hasegawa K., Watanabe Y., U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the European Bioinformatics Institute
                           NERVOUS SYSTEM, DISTINGUISHES ASTROCYTES FROM OTHER GLIA SIMILARITY: BELONGS TO THE INTERMEDIATE FILAMENT FAMILY.
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01-PEB-1990 (Rel. 1)
01-FEB-1994 (Rel. 2)
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Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;

Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;

Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
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HS11_ARATH
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DOMAIN
        modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                      This SWISS-PROT entry is copyright. It is produced through a continuous the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
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                                                                                                                                                                                                  Arabidopsis thaliana."
                                                                                                                                                                                                                                    STRAIN=CV. COLUMBIA;
MEDLINE=90158498; PubMed=2482931;
                                                                                                                                                                                                                                                                                   NCBI_TaxID=3702;
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                                                                                                                 SIMILARITY: PLANTS HAVE FOUR CLASSES OF HSP20: CLASS I AND II WHICH ARE CYTOPLASMIC; CLASS III WHICH IS IN THE CHLOROPLAST AND CLASS IV WHICH IS IN THE ENDOMEMBRANE. THIS PROTEIN BELOW
                                                                                                                                                               SUBCELLULAR SIMILARITY:
                                                                                                         TO CLASS
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169 373
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221 226
227 248
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253 373
                                                                                                                                                                                                                                                                                              Brassicaceae; Arabidopsis
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                                                                                                                                                              219:365-372(1989).
LOCATION: CYTOPLASMIC.
BELONGS TO THE SMALL HEAT
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28,
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                                             institutions as long
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SHOCK PROTEIN (HSP 17:4)
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LINKER 12.
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COIL 1A.
LINKER 1.
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LINKER 2.
COIL 2B.
; 6891CBBBF2F59AD4 CRC64;
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Pred. No. 24;
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                                            as its content
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Query Match Best Local S Matches 7

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Score 35; Pred. No.

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Best Local
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01-JAN-1990 (Rel. 13, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
17.6 KDA CLASS I HEAT SHOCK PROTEIN (HSP 17)
                            InterPro; IPR002068; -.
Pfam; PF00011; HSP20; 1.
PROSITE; PS01031; HSP20; 1.
                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II;
Brassicales; Brassicaceae; Arabidopsis
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Heat shock; Multigene family.
SEQUENCE 156 AA; 17454 MW;
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PIR; JQ0351; JQ0351. .
InterPro; IPR002068; -.
Heat shock; SEQUENCE 1
                                                                           PIR;
                                                                                                                     or send
                                                                                                                                    entities requires a
                                                                                                                                               use by non-profit institutions as long modified and this statement is not removed
                                                                                                                                                                                                                                                     -II- SIMILARITY: PLANTS HAVE FOUR CLASSES OF HSP20: CLASS I AND II WHICH ARE CYTOPLASMIC; CLASS III WHICH IS IN THE CHLOROPLAST AND CLASS IV WHICH IS IN THE ENDOMEMBRANE. THIS PROTEIN BELONGS
                                                                                                                                                                                                                                                                                                                                                Nucleic
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"An Arabidopsis thaliana cDNA clone encoding
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SUBCELLULAR LOCATION: CYTOP
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                                                                                                                     an email to license@isb-sib.ch).
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 Multigene family.
156 AA; 17547 MW;
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                                                                                                                                    license agreement (See http://www.isb-sib.ch/announce/
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 57BA4B7F1BD637DC CRC64;
                                                                                                                                                                                                                                                                                                                  SMALL HEAT
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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13.062 Million cell updates/sec
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O82741 arabidopsis
Q9skd9 arabidopsis
Q9xid4 arabidopsis
Q91qe8 arabidopsis
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O59904 kluyveromyc
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Q9zas6 streptococc
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Q9nf41 drosophila
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Q9vxp2 drosophila	Q91qe3 arabidopsis	Q9n5d4 caenorhabdi	Q9pah0 xylella fas	Q9xb56 erwinia car	Q9smf2 helianthus	P93440 oryza sativ		Q9zp84 quercus sub	Q9zs24 castanea sa	P93439 oryza sativ	Q91dg9 q heat shoc	Q9m408 quercus sub	Q9m409 quercus sub	Q9m410 quercus sub	Q9m411 quercus sub	Q9m416 quercus sub	Q9m417 quercus sub	Q9m419 quercus sub		Q9m412 quercus sub	O16103 drosophila	Q9v8k9 drosophila	Q9v6k1 drosophila	076251 drosophila	017088 caenorhabdi

## ALIGNMENTS

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RESULT
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                                                                                                                              Query Match 100.0%; Score 61; DB 4; Best Local Similarity 100.0%; Pred. No. 0.0031; Matches 12; Conservative 0; Mismatches 0
Q92495
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J. Clin. Invest. 90:2102-2109(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q92663
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Porges A.J., Redecha
Kimberly R.P.;
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01-FEB-1997 (TrEMBLrel.
01-OCT-2000 (TrEMBLrel.
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HSSP; P12319; 1ALT.
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                                                                                                                                                                                                                                                                                                                        "Novel
                                                                                                                                                                                                                                                                                                                                                                                  FISSUE=BLOOD;
                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9606;
                                                                       265 EDGNVLKRSPEL 276
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PRELIMINARY;
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cha P.B., Doebele R.,
                                                                                                                                                                                                     ; 1.
42632 MW; D33D59398CEEA699 CRC64;
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Last sequence update)
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PRT;
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375 AA.
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                                                                                                                                                          Length 374;
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RESULT
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Best Local
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Q9MZTO;
01-OCT-2000 (TrEMBLrel. J
01-OCT-2000 (TrEMBLrel. J
01-OCT-2000 (TrEMBLrel. J
                                             Yan Y., L
        Yan Y., Li X., Wang A., Zha "Molecular cloning and iden high affinity Fc receptor f Vet. Immunol. Immunopathol. EMBL; AF162866; AAF80477.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         092495;
01-FEB-1997 (TrEMBLrel. 02
01-FEB-1997 (TrEMBLrel. 02
01-0CT-2000 (TrEMBLrel. 15
FC GAMMA RECEPTOR TYPE I ()
  Receptor.
                                                                                 Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutel
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
                                                                      Bovidae; Bovinae;
NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                              structure.
                                                                                                                                                                                                                                                                                                                                                                           EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; M91555; AAA58414
                                                                                                                                                                                                                                                                                                                                                                                                                                              gamma receptor gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (DEC-1992) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human)
Eukaryota; Metazoa; (
Mammalia; Eutheria; )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Benech P.D.,
                                                                                                                                                                                                         266
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
                                                                                                                  GAMMA RECEPTOR I
                                                                                                                                                                                                                   1 EDGNVLKRSPEL
                                                                                                                                                                                                                                                                                                                                      M91553; AAA58414.1; J
M91554; AAA58414.1; J
S45709; AAD13842.1; J
S45707; AAD13842.1; J
S45708; AAD13842.1; J
S45704; AAD13842.1; J
                                                                                                                                                                                                        EDGNVLKRSPEL
                                                                                                                                                                                                                                                                                                      PF00047;
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                                                                                                                                                                                                                                                                                                                                                                                            м91552;
                                                                                                                                                                                                                                                                                                                                                                                                     м91551;
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12; Conser
                                                   FROM N.A.
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                                                                                                                                                                                                                                         Conservative
                                                                                                                                                             PRELIMINARY;
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AAA58414.1;
AAA58414.1;
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               , Zhang G.;
identification of full-length
for bovine IgG (FC gamma R
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C GAMMA RIB=FC GAMMA REC
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Pred.
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Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                  A84D464C70DD0F91 CRC64;
                                                                                                                                                                                                                                        Mismatches
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No.
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                                                                                             Euteleostomi;
                              cDNA encoding
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                                                                                    Bovoidea;
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RA Fosler C. Gabrielian A.E., Garg N.S., Gelbart M.M., Glasser K., Ra Glodek A., Gong F., Gorrell J.H., Gu'Z., Guan P., Harris M., RA Glodek A., Gong F., Gorrell J.H., Gu'Z., Guan P., Harris M., RA Glodek A., Gong F., Gorrell J.H., Gu'Z., Guan P., Harris M., RA Glodek A., Gong F., Gorrell J.H., Gu'Z., Guan P., Harris M., RA Harris D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C., RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X., RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D., Lai Z., RA Lasko P., Lei Y., Levitsky A.A., Li J., Li Z., Liang Y., Lin X., RA McIntosh T.C., McLeod M.P., McPherson D., L., McPherson D., McPher
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Best Local S
Matches 9
  Borkova D., Minana B., "Sequencing the distal Submitted (FEB-1999) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ballew R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M.,
Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
Dodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P.
Durbin K.J., Evangelista C.C., Ferriera S., Fleischmann W.
                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-1999
01-MAY-1999
01-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D., Wan K.H., Doyle C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.
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Amanatides P.G., Scherer S.E., Li P.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=20196006; PubMed=10731132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pterygota; Ne Ephydroidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota;
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9; Conserv
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Metazoa; Arthropoda;
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81.8%;
                   Kafatos F.C.;
X chromosome of
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comosome of Drosophila melanogaster.";
EMBL/GenBank/DDBJ databases.
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Pred. No. 0.
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Galle R.
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Smith H.O.,
                                                                                                                                                                                                                                                                                                                                                                     D.L.,
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01-NOV-1996
01-JUN-2000
                                                                                             EMBL; U19
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (FEB-1999) to the EEMBL; AE003426; AAF45849.1; --
EMBL; AL035395; CAB37612.1; --
FLYBASE; FBB00000644; Fcp3C.
SEQUENCE 99 AA; 11150 MW;
                                                                                                                                                                                                                                                                                                                                             Underwood A.P., Urrestarazu L.A., Vandenbol M., Verhasselt F., Verens B., Volckaert G., Voss H., Wambutt R., Wedler H., Zimmermann F.K., Zollner A., Hani T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycetes;
                                                                                                                        Submitted
                                                                                                                                    Cherry J.M.;
                                                                                                                                                SEQUENCE FROM N.A.
STRAIN~S288C (AB972);
                                                                                                                                                                                                                                 SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. STRAIN=S288C (AB972);
                                                                                                                                                                                                                                                                                                                                                                                              Muller-Auer S., Nentwich U., Portetelle D., Purnelle B., F Scharfe M., Scherens B., Scho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=97313267; PubMed-
Johnston M., Hillier L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CHROMOSOME XII COSMID
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                                                                                                                                                                                                       Waterston
                                                                                                                                                                                                               STRAIN-S288C (AB972);
                                                                                                                                                                                                                                                           Submitted (JAN-1995)
                                                                                                                                                                                                                                                                                                                             Nature
                                                                                                                                                                                                                                                                                                                                        "The nucleotide sequence of Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                    Benes V., Bruckner M., Delius H., Dubois E., Dusterhoft A.,
Entian K.D., Floeth M., Goffeau A., Hebling U., Heumann K.,
Heuss-Neitzel D., Hilbert H., Hilger F., Kleine K., Kotter P.,
Louis E.J., Messenguy F., Mewes H.W., Miosga T., Mostl D.,
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 1 EDGNVLKRSPEL
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                          Similarity
8; Conserv
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                            Conservative
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Pred. No.
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                         ced. No. 2.7;
Mismatches
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Best Local
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Q39012;
01-NOV-1996
                                                      SEQUENCE FROM N.A.
STRAIN-CV. COLUMBIA;
Liu S.X., Vaysberg M., Etgu P.
                                                                                                                                                                                                                                                                                                            01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-CT-2000 (TrEMBLrel. 15, Last annotation update)
SHAGGY-LIKE PROTEIN KINASE IOTA (TZDZ3.9 PROTEIN)
ASK IOTA OR GSK1 OR TZDZ3.9.
ARABIODPSIS thaliana (Mouse-ear cress).
                                                                                                                                                                                                                                                                        Eukaryota; Viridiplantae; Embryophyta; Trache
Magnoliophyta; eudicotyledons; core eudicots;
Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (OCT-1999) to the EMBL/GenBank/DDBJ databases EMBL; AL121805; CAB58072.1; SEQUENCE 299 AA; 32007 MW; C7224AFCDC0809B6 CRC64;
                                   Toriumi M., Yu G., Ch
Altafi H., Brooks S.,
                                                                                                                                                                                           SEQUENCE FROM N.A.
TISSUE=ROOTS;
DOTRICAL SCHWebel-Dugue N., Thomas M., Lecharny A., Kreis
Plant Physiol. 113:306-306(1997).
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         Hansen N.F., Johnson-Hopson C., Nguyen M., Palm C.J., Shinn P.,
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Sequencing
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01-OCT-2000
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                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cadieu, Dreano, Lelaure, Galibert F.; ing the distal X chromosome of Drosophila melanogaster."; d (OCT-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                  (AUG-1997)
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                                               Chan
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Endopterygota; Diptera; Brachycera; Muscomorpha;
Llidae; Drosophila
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                                   an A., Chung
Buehler E.,
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Pred. No.
          Southwick A.,
                                                                                                                                        Lim J.H.,
                                                            Lee J.M.,
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                M., Goldsmith A., Liu A., Smith A Chao Q., Conn L., Conway A.B., han S., Kim C., Lam B., Mirana.
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licots; Rosidae; eurosids II;
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             Davis
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                        Lam B., Miranda M.,
                                              Smith A.,
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01-NOV-1999 (TrembLrel.
01-OCT-2000 (TrembLrel.
651AA LONG HYPOTHETICAL
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TIGR; VC0302; -.
SEQUENCE 396 A
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Dodson R.J., Haft D.H., Hickey E.K., Peterson J.D., Umayam
Gill S.R., Nelson K.E., Read T.D., Tettelin H., Richardson
Ermolaeva M.D., Vamathevan J., Bass S., Qin H., Dragoi I.,
McDonald L., Utterback T., Fleishmann R.D., Nierman W.C., W
Salzberg S.L., Smith H.O., Colwell R.R., Mekalanos J.J., Ve
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STRAIN-EL TOR N16961 / SEROTYPE 01

MEDLINE-20406833; PubMed-10952301.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fraser C.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria; Proteobacteria;
NCBI_TaxID=666;
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PROSITE; PS00107; PROTEIN_KINASE_ATP; 1.

PROSITE; PS00108; PROTEIN_KINASE_DOM; 1.

PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.

SEQUENCE 407 AA; 46024 MW; 96BC4C537
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'DNA sequence of both chromosomes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vibrio cholerae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AF019927; AAB71545.1;
EMBL; AC068143; AAF82167.1;
HSSP; P24941; 1JSU.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MENDEL; 13957; Arath;2319;13957.
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                                                                                                                                                                                                                                               1 EDGNVLKRSPE 11
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                                                                                                                                                                                                                      QDGNALKAEPE 393
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence of BAC T2D23 from Arabidopsis thaliana chromosome 1."; itted (UUL-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                         Similarity 7; Conserv
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                                                                                                            PRELIMINARY;
                                                                                                                                                                                                                                                                                                       Conservative
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(TremBLrel. 15, Last sequence update)
(TremBLrel. 15, Last annotation update)
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72.7%;
12, Created)
12, Last sequence update)
15, Last annotation update)
LONG-CHAIN-FATTY-ACID--COA
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Pred.
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Pred. No. 11;
1; Mismatches
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ed. No. 25;
Mismatches
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yam L.A.,
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Conservative

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Gaps

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Score 38.5; Pred. No. 6

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Length Indels

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EGNITHSTLLKRSPEL

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Best Local S
Matches 7
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01-MAY-1999
01-MAY-1999
01-MAY-2000
   SEQUENCE
                                                                 Podbielski A., Leonard B.A.,
"The group A streptococcal c
uptake of essential amino ac
                   PFAM; PF00496; SBP_bac_5;
                                     Mol. Microbiol. 28:1323-1334(1998).
EMBL; U78968; AAC67217.1; -.
                                                          protease."
                                                                                             MEDLINE=98343814; PubMed=9680220;
                                                                                                      STRAIN=CS10
           Lipoprotein
                              INTERPRO; IPRO00914;
                                                                                                                                   NCBI_TaxID=1314;
                                                                                                                                             Streptococcus
                                                                                                                                                      Bacteria; Firmicutes;
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                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                 432
                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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Nomura N., Sako Y., Kikuchi H.;
"Complete genome socciety."
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Jin-no K., Takahashi M., Sekine M., Baba S., An
Hosoyama A., Fukui S., Nagai Y., Nishijima K.,
Takamiya M., Masuda S., Funahashi T., Tanaka
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NCBI_TaxID=56636;
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Archaea; Crenarc
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61145 MW;
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on update)
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                                                               expression
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01-MAY-2000
01-MAY-2000
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01-OCT-2000
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STRAIN-CV. CASTELLANA; TISSUE-ETIOLATED
STRAIN-CV. F. RSTEBAN R., DOPICO B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cicer arietinum (Chickpea) (Garbanzo).

Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Magnoliophyta; Liliopsida; NCBI_TaxID=4577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2000 (TrEMBLrel. 14, I
PUTATIVE ENOYL COA HYDRATASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Jittery, a low-copy, Mu-related BSMV infection.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TRANSPOSASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q9SMK7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yan X., Li J., Cowperthwaite M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zea mays (Maize)
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                                                                                                                                                                                                                                                                    586
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                                                                                                                                                                                                                                                                                                                               1 EDGNVLKRSPE 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 EDGNVLKRSPEL
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                                                                                                                                                                                                                                                                 DDGNLLEEKPE 596
                                                                                                                                                                                                                                                                                                                                                                                                Similarity 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 7; Conserv
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(TrEMBLrel.
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                                                                                                PRELIMINARY;
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(TrEMBLrel. 14, Last annotation update)
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58.3%;
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Created)
Last sequence update)
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Last annotation update)
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                                                                                                   PRT;
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1.1e+02;
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Best Local (
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Best Local (
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Q53734;
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Theme and Kluyveromyces lac cerevisiae and Kluyveromyces lac genetics 148:1021-1029(1998).
EMBL; AF035007; AAC39438.1; -
SFOHENCE 1314 AA; 146734 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Fuńgi; As
Saccharomycetaceae;
NCBI_TaxID=28985;
                                                    01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-CCT-2000 (TrEMBLrel. 15, Last annotation update)
HYPOTHETICAL 30.1 KDA PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Structural analysis of plasmid pSN2 in Staphylococcus aureus: involvement in enterotoxin B production.";
J. Bacteriol. 149:642-649(1982).
EMBL; J01763; AAA91889.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. MEDLINE-82119973; Pubmed-7056699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Staphylococcus aureus
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
PLASMID PSN2 INCLUDES AN UNKNOWN GENE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kluyveromyces lactis.
Eukaryota; Fungi; Ascomycota;
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SILENT INFORMATION REGULATOR
Arabidopsis thaliana (Mouse-ear cress)
Eukaryota; Viridiplantae; Embryophyta;
                                     F7H19.100 OR AT4G22920
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Khan S.A., Novick R.P
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Pred. No.
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Pred. No. 27;
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                                                                                                                                              PRT;
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                                                                                                                                                                                                                                                                                                                                       Length 158;
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Search completed: June 4, 2001, 12:23:30 Job time: 593 sec
                                                                                                                          망
                                                                                                                                                                                                              Query Match
Best Local Similarity 63.6
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
BU Arabidopsis sequencing project;
BU Arabidopsis sequencing project;
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AL031018; CAA19807.1;
EMBL; AL161558; CAB79247.1;
Hypothetical protein.
Hypothetical protein.
BEQUENCE 268 AA; 30052 MW; 1B3F128F18EA79B2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.

Peters S.A., van Staveren M., Dirkse W., Stiekema W., Bancroft I.,

Mewes H.W., Mayer K.F.X., Schueller C., Bevan M.;

Submitted (JUL-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II;
Brassicales; Brassicaceae; Arabidopsis.
NCBI_TaxID-3702;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
Peters S.A., van Staveren M., Dirkse W., Stiekema W., Mewes H.W.,
Lemcke K., Mayer K.F.X.;
Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                     2 DGNVLKRSPEL 12
|||:| |||
170 DGNLLNNYPEL 180
                                                                                                                                                                                                                                               60.7%;
                                                                                                                                                                                                                     Score 37; DB 10; pred. No. 60; 1; Mismatches 3
                                                                                                                                                                                                                          3; Indels
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                        Score
   Match
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'SIDS6/gcgdata
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                                                                                                                                                        344
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                                                                                                                                                                                                          DВ
                                W60560
R91439
Y96183
R91438
W00859
W80448
W97833
W97833
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                                                                                                                                                                                                                                                          SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        geneseqp/AA1988.DAT:
                                                                                                                                                                                                        Description
                                                  Human
                                                                                                                                                                    Oligopeptide from
                                                                    Human
                                                                                    Human
                                                                                                                      Human
                                                                                                                                      Human
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n FcRI (cDNA c
n FcRI (cDNA c
high affinit
cancer assoc
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	P8098	R76	20	R76	R56		W3(					_	_	_		_	۰					G2602		R6	R67	R87	R1327	₩	R1327	W8836	R2255	R2081	R2254	0 W86195
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	Sequence of fusion	5-11k	in kinase (C	HRR25-lik	in kina	Amino acid sequenc	V aden	<ul> <li>Cardiac adenylyl c</li> </ul>	c adenylyl	pe V adenylyl	rreus LovC	Mal prote	acid	Mouse BH3 interact	BID trunca	an transcripti	tein involved	mays protein	mays	protein	mays protein	mays prote	protei	male	Φ.	noqlobu	ocyte Activa	Human lymphocyte a	edrin-solubl	lymphocyte	macrophage-	macrophage-	n macrophage	Human Fc receptor

## ALIGNMENTS

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W60560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                  Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; detecell transformation; mutation; anti; oncogene.
  De Kruif CA,
                                                                                08-OCT-1996;
                                                                                                                        07-OCT-1997;
                                                                                                                                                                                                      W09815833-A1.
                                                                                                                                                                                                                                                                                                                                                                  Oligopeptide from extracellular domain of CD64
                                                                                                                                                                                                                                                                                                                                                                                                          18-AUG-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                  W60560;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          W60560 standard;
                                        (UYUT-) RIJKSUNIV UTRECHT.
                                                                                                                                                                L6-APR-1998
Logtenberg T;
                                                                                96EP-0202791.
                                                                                                                        97WO-NL00557.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide; 12 AA.
                                                                                                                                                                                                                                                                                                         immunofluorescent analysis; detection;
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B43683

Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying specific antibodies

WPI; 1998-240964/21.

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RESULT
R91439
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Best Local Similarity
Matches 12; Conser
                                                                                                                              01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               forms of the same protein, including bispecific antibodies that bind to two non-overlapping epitopes on the same monomeric antigen or two epitopes on different molecules. The genes/oligonucleotides that encode selected peptides can be isolated and used for recombinant production of antibodies. Selected antibodies are used, e.g. for immunohistochemical or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                surface of a replicable display package, synthesising oligopeptides derived from the target protein on a solid phase, and testing for binding between the peptide and oligopeptides. The method is used to screen large peptide libraries, especially to detect antibodies, or their fragments, that bind to cell markers or that can differentiate between different forms of the same protein, including bispecific antibodies that bind to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            method for the identification of a peptide able to bind specifically to a target protein. The method comprises displaying the peptide on the surface of a replicable display backage surface of a replicable display backage surface.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic oligopeptides W60537-61 are derived from the extrace domain of CD64. They were synthesised on 25 polyethylene rods adjacent, non-overlapping sets. Peptide specific phages were infinity binding to the CD64-covered rods. The specification of the CD64-covered rods.
N-PSDB; T14718
                                               Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                         Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FcRI; Fc receptor.
               WPI; 1996-200279/20
                                                                                                                                                                                                                 25-FEB-1988;
                                                                                                                                                                                                                                                 09-APR-1996.
                                                                                                                                                                                                                                                                             US5506126-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R91439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R91439 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antibodies. Selected antibodies are used, e.g. immunofluorescent analysis, and also to detect
                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    by mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 29; 40pp;
                                                                               (GEHO ) GEN HOSPITAL CORP
                                                                                                                18-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 ELQVLGLQLPTP 12
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1 elqvlglqlptp 12
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                                                                                                                              92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
                                                                                                               9308-0139273
                                                                                                                                                                                                                 88US-0160416
                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                             /note= "amino acid 58 is Leu in clone p135
    translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 60;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.00013;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        immunotherapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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RESULT
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA: clones, pl35 (T14717) and p90 (T14719), coded for a variants of the FCRI sequence; the C-terminal sequence of the p98 product is truncated compared with those of the p135 and p90 products (see also R91438 and W00859). A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
 Misc-difference
                             Misc-difference
                                                                                   Misc-difference
                                                                                                                                             Misc-difference
                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                          panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy;
immune disorder; infection; asthma; immune-complex disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Y96183 standard; Protein; 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The amino acid sequence (R91439) cDNA clone, p98 (T14718), obtd. immunoselection cloning method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins % \left( 1\right) =\left\{ 1\right\} 
                                                       Misc-difference
                                                                                                               Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Macrophage; FCRI; cell surface antigen; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human macrophage-specific FcRI.
                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
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|elqvlglqlptp-288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           multiple sclerosis; parasitic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                          /note=
                                                                                                                                                                                                                                                      /note=
/note=
126
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117
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123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FCRI is a high affinity receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of human FcRI
                                                                                                  GAA"
                ACC.
                                                                                                                                                           CAG"
                                            TTC"
                                                                       GCA"
                                                                                                                                                                                                                  TCC"
                                                                                                                                                                                                                                                                                                    cgg."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ĎВ 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       immunoselection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 344;
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The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
                                                                                                                physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                   immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the FC portion of ISC normally located on the cell surfaces of macrophages. The second surface of the cell surfaces of macrophages.
                                                           treatment of immune-mediated infections, diseases, and diso
animals, including humans. These disorders include asthma,
                                                                                       proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis an
                                                                                                                                                         based upon transient expression of a CSA in eukaryotic
                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                           Isolated nucleic acid molecule encoding the CD19 cell surface useful for immunodiagnosis and immunotherapy of immune-mediate infections or disorders, e.g. asthma, immune-complex disease,
                                                                                                                                                                                                                                                                                                                                                                         Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN
                                                                                                                                                                                                                                                                                                                                                                                                                                           13-JUL-1989;
23-MAR-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992;
25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-OCT-1998;
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           interfere with such bonding, or
                                                                                                                                                                                                                                                                                                                                                                                                   HOSPITAL CORP
                                                                                                                                                                                                                                        Column 53-55;
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89US-0379076.
90US-0498809.
90US-0553759.
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           cause it to occur on
                                                                                                                                                                                                                                                                         immune-mediated nplex disease, parasitic
                                                                          and disorders in
                                                                                                                                                                      The method is
                                                                                                                                                        cells and
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Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                            25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                (T14718), coded for an FCRI (R91439) having a different C-terminal sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein of FCRI and a receptor ligand will the potencies of antibodies in therapy.
                                                            for the Fc portion of IgG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FcRI sequence, and a third clone,
                                                                                                                  cDNA clone,
                                                                                                                                                    Example 10; Column 55-56; 79pp;
                                                                                                                                                                               Cloning of cDNA encoding cell surface antigen of diagnostic and therapeutic proteins
                                                                                                                                                                                                                      N-PSDB;
                                                                                                                                                                                                                                                          Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                 01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                    09-APR-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cell surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human FcRI (cDNA clone p135 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R91438;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R91438 standard;
                                                                                                                                                                                                                                                                                    (GEHO ) GEN HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                          25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                  immunoselection cloning method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    277 elqvlglqlptp 288
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                                                                                                               amino acid sequence (R91438)
A clone, p135 (T14717), obtd.
                                                                                                                                                                                                                      1996-200279/20
DB; T14717.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ELQVLGLQLPTP
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374 AA
                                                                                                                                                                                                                                                            Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      344 AA;
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                                                                                                                                                                                                                                                                                                                          92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
                                                                                                                                                                                                                                                                                                               93US-0139273
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vector;
                                                                                                                                                                                                                                                                                    CORP.
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translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "amino acid
translated
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                                                                                                              obtd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FcRI; Fc receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 60;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ā
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunoselection;
                                                                                                                                                    English
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                                                                                                   FcRI is a high affinity receptor
                                                                                                                             of human FcRI was detd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 25 is Thr
1 products"
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Sequence

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Matches 12
                                                                                                                                                                             The amino acid sequence (W00859) of human FCRI was detd. from a cDNA clone, p90 (T14719), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the FC portion of IgG, normally located on cell surfaces of macrophages. Other cDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of
                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
                                                                                                                                            Sequence
                                                                                                                                                                 potency
                                                                                                                                                                                                                                                               Example 10;
                                                                                                                                                                                                                                                                          Cloning of cDNA encoding cell surface antigen - useful for isolation of diagnostic and therapeutic proteins
                                                 277
                                                                                                                                                                                                                                                                                                                                                   Aruffo
                                                                                                                                                                                                                                                                                                                                                                      (GEHO)
                                                                                                                                                                                                                                                                                                                                                                                            18-OCT-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US5506126-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cell surface antigen; cloning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human FcRI (cDNA clone p90 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W00859;
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                                                          1 ELQVLGLQLPTP 12
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111111111111
277 elqvlglqlptp 288
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DB; T14719.
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                                                                                       1 Similarity 100.0%;
1 Similarity 100.0%;
12; Conservative
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of antibodié
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12; Conserva
                                                                                                                                            374
                                                                                                                                                                                                                                                          Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                      HOSPITAL CORP
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                                                                                                                                            AA;
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88US-0160416.
89US-0379076.
90US-0553759.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vector; FcRI;
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                                                                                    Score 60; DB Pred. No. 0.0); Mismatches
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Pred. No. 0.0054;
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                                                                                                0.0054;
                                                                                                          DВ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunotherapy;
                                                                                                        Length
                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 374;
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                                                                         WPI; 1998-609251/51.
N-PSDB; V63456.
                                                                                                                                                      Aruffo A,
                                                                                                                                                                          (GEHO ) GEN HOSPĮTAL CORP.
                                                                                                                                                                                                                                            01-DEC-1992;
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New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences

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900S-0553759 970S-0861205 88US-0160416 89US-0379076 92US-0983647 97us-0861205

Example 10; Column 53-54; 75pp; English.

This is the amino acid sequence of human Fc receptor I (FCRI), as deduced from cDNA clone pl35 (see V63456) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2

RESULT W80448

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Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                efficiency expression vectors (see V63441 and V63444) which allow
the generation of very large mammalian expression libraries. The
purified genes and proteins are useful for immunodiagnostic and
immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocyttes (see W80440-55). The isolated genes can be expressed in a prokaryotic or enkaryotic host cells to produce the encoded protein. The invention also provides high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adhesion to an antibody-coated substrate.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          eukaryotic cells and selection of cells expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             also obtained
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             374 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immune-mediated infections, diseases,
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- 85
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171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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Pred. No. 0.0054;
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25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             were also obtained. A novel method for cioning curves come expression libraries is based on transient expression of an antigen in enkaryotic cells and selection of cells expressing the antigen by the control of the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method is useful for the method
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                                                                                      07-JUN-1999
                                                                                                                                                                             W97834;
                                                                                                                                                                                                                                                               W97834 standard; Protein; 374 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
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N-PSDB; X07372.
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Human Fc receptor I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      including humans
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ilarity 100.0%;
Conservative 0
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89US-0379076
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Clones pl35 (see V63456) and p98 (see X07373)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   method for cloning cDNAs from mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 60; DB 19;
Pred. No. 0.0054;
0; Mismatches 0;
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Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;

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cloning.

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from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expression the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-63) encoding cell surface antigens from mammalian lymphocytes (see W80440-55). The isolated
                                                                                                                                                                                                                                          New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                 Example 10; Column 53-54; 75pp; English.
                                                                                                                                                         This is the amino acid sequence of human Fc receptor I (FCRI), deduced from cDNA clone p98/X2 (see X07373) isolated using a rimmunoselection cloning method from a cDNA library expressed in
                                                                                                                                                                                                                                                                                                                                                 Aruffo A,
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Misc-difference 171
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                                                                                                                                          The cDNA library was constructed from polyA RNA of cells
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90US-0553759.
97US-0861205.
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Misc-difference 55
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                                                                                                                                                                                                                                                                                                                                                                                                   macrophage; rckl; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Macrophage; FcRI; cell surface antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 produce the encoded protein. The invention also provides high efficiency expression vectors (see v63441 and v6344) which allow the generation of very large mammalian expression libraries. The immunotherapeutic applications, including the diagnosts and treatment of immune-mediated infections, diseases, and disorders of animals, including humans.
                                                                                                                                                                                                                                                                                                                                                    Misc-difference
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ne encoded protein. The invention also provides h
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Misc-difference

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physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coates of the coate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone p135 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasitic
surfaces other than macrophages, is useful protein of FCRI and a receptor ligand will the potencies of antibodies in therapy.
                                   immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 10; Column 53-55; 75pp; English
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25-FEB-1988;
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305
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    note= "encoded by AAT'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "encoded by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "encoded by AAT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "encoded by TTC"
                                                                                                                                                                                                                                                                                                     These disorders include asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTG"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAA'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCA
                                                                                                                                                                                                                                                                                                                                                  and disorders in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The method is
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Matches 12; Conser

Conservative

100.0%; Score 60; 100.0%; Pred. No. tive 0; Mismatc

Mismatches

0.0054;

Length 374; Indels

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1 ELQVLGLQLPTP 12 elqvlglqlptp 288

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Matches 12; Conserv
                                                                                                                                                                                                                                 receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical compositions containing recombinant proteins may be used to treat or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neutrophil; eosinophil; HIV; IGG; immunosuppressive; antirhematic; cytostatic; antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                       neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FegammaRI is also known as CD64. Fer receptors. Play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds cells expressing Fer receptors. FegammaRI molecules are expressed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          monocytes and macrophages, but expression can also be induced on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-367968/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-JUN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EP1006183-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               systemic lupus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human high affinity Fc receptor, FcgammaRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Y96226;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Y96226 standard; Protein; 374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; A27466
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                        rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10
                                                                                                                                 374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence is the human high affinity Fc receptor,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 26-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               erythematosus; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98EP-0122969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98EP-0122969.
                                                                                                                                                                                                                 diseases,
                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             60pp; English.
                                                                                                                                                                                        X
                                                                                                                                                                                                                 allergies
                           Score 60; DB 21; Pred. No. 0.0054;
                                                                                                                                                                                                                 or tumours,
                                                    DB 21;
                                                    Length 374;
                                                                                                                                                                                                                 especially AIDS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FcgammaRI
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                                                                                                                                                                                                          vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and polypeptides can be used for preventing, treating or ameliorating medical conditions and diagnosing pathological conditions. Polynucleotides, polypeptides, antibodies, agonists and antagonists from the present invention may be used to treat immune disorders by activating or inhibiting the proliferation, differentiation or mobilisation of immune cells, to treat disorders of haematopoietic cells, autoimmune disorders, modulate reactions, graft versus host disease and organ rejection, modulate haemostatic or thrombolytic activity, modulate inflammation, cancers, cardiovascular disorders, neurological disease and bacterial or viral infections. The peptides, nucleotides, antibodies, agonists and antagonists may be also be used in drug screens. C78449 to C78457 and 194420 represent sequences used in the exemplification of the present invention.
                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C77607 to C78448 encode the human cancer associated proteins given in B43398 to B44239. The proteins can have activities based on the tissues and cells the genes are expressed in. Example of activities include: cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antithyroid; antiallergic; antibacterial; antiviral; dermatological;
neuroprotective; cardiant; thrombolytic; coagulant; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 11; Page 1739-1740; 2352pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vasotropic; antipsoriatic and antiangiogenic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel isolated nucleic acids comprising sequences encoding useful for treating or diagnosing e.g. cancer
                                                                                                                                                                                                  Invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosen CA, Ruben SM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-MAR-2000; 2000WO-US05882
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vasotropic; antipsoriatic; antiangiogenic; gene therapy; inflammati immune disorder; haematopoietto cell disorder; autoimmune disorder; altergic reaction; graft versus host disease; organ rejection; haemostatic; thrombolytic; cardiovascular disorder; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           diagnosis; cytostatic; proliferative; vulnerary; immunomodulator; antidiabetic; antiasthmatic; antirheumatic; antiarthritic; antiviral; antiinflammatory; antithyroid; antiallergic; antibacterial; cardiant; dermatological; neuroprotective; thrombolytic; coagulant; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B43683;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B43683 standard; Protein; 399 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; cancer associated gene; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human cancer associated protein sequence SEQ ID NO:1128
                                      Local
           l Similarity
.12; Conserv
                                                                                                                                          399 AA;
        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease; drug screening.
                                 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cardiovascular
  Score 60; DB 21;
Pred. No. 0.0057;
); Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disorder; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       detection; cancer;
                                                    Length 399;
  Indels
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ELQVLGLQLPTP 12 

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Best Local Query Match Matches

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Conservative

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86.7%; 91.7%;

Score 52; DB Pred. No. 0.15 0; Mismatches

0.15; DB 20;

Length 377; Indels

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                                                                                                                                                      v81213) were isolated using a rapta method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extreorporeal interleukin-2 induction therapy. DNA sequence analysis revealed that the cDNAs encoded therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-MAR-1990;
13-JUL-1990;
07-JUN-1995;
                       and molecular cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. CD40 cDNA (see V81198) is specifically claimed.
                                                                       type I integral membrane proteins with 3 extracellular immunoglobulin domains. The invention provides a novel method for cloning cDNAs from mammalian expression libraries. This is based on transient expression of an antigen in eukaryotic cells and physical selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation
                                                                                                                                                                                                                          This polypeptide human Fc receptor I (FcRI). FcRI cDNAs (see V81213) were isolated using a rapid immunoselection cloning
                                                                                                                                                                                                                                                                                                                        ODNA encoding human CD40 antigen - useful for cloning cDNA encoding cell surface antigens, constructing cDNA libraries, expression vectors for expression in eukaryotic cells or their fragments
                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-069813/06
N-PSDB; V81213.
                                                                                                                                                                                                                                                                                                   Example 10; Column 55-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oquendo C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Allen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US5849898-A
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13-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human Fc receptor I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W86195;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amiot M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92US-0983647.
88US-0160416.
89US-0379076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CD36; cell surface antigen; human; cDNA library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          900S-0498809
900S-0553759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-0485447
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "this sequence contains a considerable number of differences from the sequence deduced from the DNA sequence given in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                Aruffo A, C
                                                                                                                                                                                                                                                                                              79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the specification'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      377
                                                                                                                                                                                                                                                                                                                                                                                                                                               Camerini D, I
, Stamenkovic
                                                                                                                                                                                                                                                                                                                                                                                                                                                Lauffer L;
c I, Stengelin
                                                                                                                                                                                                                                                               FCRI cDNAs (see
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                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                      This amino acid sequence was predicted from the cDNA sequence of cDNA clone p98/X2. It differs from the sequence predicted from clone p135 (see Q21178) at position 25; p135 encodes a Ser residue while clone p98/X2 predicts a Thr residue. At position 58, p90 (see Q21180) predicts Val and p98/X2 predicts Leu. Sequences predicted from all 3 clones show the typical features of a type I
             R20811;
                                              R20811 standard;
                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                        integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six potential N-linked glycosylation sites and six
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 94a; 160pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rapid immunoselection cloning technique; cell surface antigen; immunodlagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human macrophage-specific FcRI receptor encoded by clone p98.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R22549;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R22549 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-1990;
                                                                                                                               277
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{|||| ||||||||
280 elqvlqlqlptp 291
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                                                                                                                                                            1 ELQVLGLQLP 10
                                                                                                                                                                                                                                                                                                                            residues distributed among three C2 set Ig-related usion protein of FCRI and a receptor ligand will be
                                                                                                                             elqvlglqlp 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aruffo A,
                                                                                                                                                                                                                                                                                                               the potency
                                                                                                                                                                                                                                                                            344
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                                              Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amiot M
                                                                                                                                                                                                           80.0%;
                                                                                                                                                                                                                                                                                                               of antibodies
                                            374 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    344
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0; Mismatches
                                                                                                                                                                                                                                                                                                           in therapy.
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0.7;
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helpful to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  distributed among three C2 set Ig-related domains of FcRI and a receptor ligand will be helpful to i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 typical features of a type I integral membrane protein and include a short hydrophobic signal sequence, a single 21-residue hydrophobic membrane-spanning domain, and a short, highly charged cytoplasmic domain. The extracellular portion contains six
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    position 25; pl35 encodes a Ser residue and the other two clones predict a Thr residue. At position 58, pl35 predicts Leu and p90 predicts Val. Sequences predicted from all 3 clones show the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This amino acid sequence was predicted from the cDNA sequence of cDNA clone p135. It differs from the sequence predicted from cDNA clones p98/X2 and p90 (see Q21179 and Q21180, respectively) at position 25; p135 encodes a Ser residue and the other two clones
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Rapid immunoselection cloning technique; immunodiagnosis; high affinity receptor.
                                                              Human macrophage-specific FcRI receptor encoded by clone p90.
                                                                                                            21-MAY-1992
                                                                                                                                                        R22550
                                                                                                                                                                                                  R22550 standard; Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             potency of antibodies in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New CD53 cell surface antigen and DNA encoding immuno-therapy and diagnosis of haematopoietic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-JAN-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              potential N-linked glycosylation sites and six Cys residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; Q21178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1992-056864/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seed B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JUL-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEHO-) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                          277 elqvlglqlp 286
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                                                                                                                                                                                                                                                                                                                                                      ELQVLGLQLP
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odiagnosis; high affinity receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Aruffo A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 94a;
                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                        (first entry)
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N-PSDB; Q21180.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seed B, Aruffo A, Amiot M;
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Copyright (c) 1993 - 2000 Compugen Ltd.
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3-794-695A-50
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	PCT-US95-02251-3	US-08-485-049D-2	US-08-485-863A-2	US-08-487-077A-2	US-08-787-739-2	US-08-485-862B-2	US-08-486-756A-2	US-08-477-504A-2	US-08-481-658B-2	US-08-839-008-9	US-08-335-469-2	US-08-787-739-87	US-08-787-739-51	.US-08-485-049D-51	US-08-485-863A-51	US-08-487-077A-51	US-08-485-862B-51	US-08-486-756A-51
	Sequence 3, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 9, Appli	Sequence 2, Appli	Sequence 87, Appl	Sequence 51, Appl	Sequence 51, Appl	Sequence 51, Appl	Sequence 51, Appl	Sequence 51, Appl	Sequence 51, Appl
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ALIGNMENTS

## US-08-416-478A-6 Sequence 6, Applica Patent No. 5773578 GENERAL INFORMATION: TITLE OF INVENTION: TITLE OF INVENTION: APPLICANT: Hercend, APPLICANT: Triebel, CORRESPONDENCE ADDRESS NUMBER OF SEQUENCES: COUNTRY: CITY: Washington STREET: ADDRESSEE: 20004 D.C. Application US/08416478A E: BROWDY AND NEIMARK, P.L.L.C. 419 Seventh Street, N.W., Suite 400 U.S.A. Hercend, Thierry New Proteins Produced By Human Lymphocytes, DNA Sequences Encoding These Proteins And Their Pharmaceutical And Biological Uses Frederic

COMPUTER READABLE FORM:	
COMPUTER: IBM PC compatible	
OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Datentin Release #1 0 Version #1 25	
H	
APPLICATION NUMBER: US/08/416,478A	
FILING DATE: 04-APR-1995	
CLASSIFICATION: 435	
PRIOR APPLICATION DATA:	
APPLICATION NUMBER: US 07/854,644	
FILING DATE: 08-SEP-1992	
ATTORNEY/AGENT INFORMATION:	
NAME: YUN, Allen C.	
REGISTRATION NUMBER: 37,971	
REFERENCE/DOCKET NUMBER: HERCEND=1A	
TELECOMMUNICATION INFORMATION:	
TELEPHONE: (202) 628-5197	
TELEFAX: (202) 737-3528	
INFORMATION FOR SEQ ID NO: 6:	
SEQUENCE CHARACTERISTICS:	
LENGTH: 380 amino acids	
TYPE: amino acid	
TOPOLOGY: linear	
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3-08-416-478A-6	

Best Local S
Matches 8

Similarity 8; Conserv

65.0%; 72.7%;

Score 39; DB. Pred. No. 13; 1; Mismatches

DB.1; Length 380; 13; 2

Indels

0;

Gaps

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Conservative

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Query Match
Best Local Similarity
Thes 8; Conserve
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                                                                                                                                                                                          Sequence 6, Application US/08394442B Patent No. 5976877
                                                                                                                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATFORNEY, Allen NAME: YUN, Allen NAME: YUN, Allen NUMBER: 37, 5, 2000 NUMBER: 47, 5, 2000 NUMBER: HERCF REFERENCE/DOCKET NUMBER: HERCF TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: 4202) 737-3528
                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (202) 737-35
NFORMATION FOR SEQ ID NO:
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                                                                                      APPLICANT:
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PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US 08/416,478
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
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                                                                                                                                                       INFORMATION:
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amino acid
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                                                                                      Triebel,
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SYSTEM: PC-DOS/MS-DOS
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                                         New Proteins Produced By Human
                                                                                                                 Thierry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.0%;
Lymphocytes, DNA Sequences Encoding These Proteins And
                                                                                      Frederic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US 07/854,644
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 39;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HERCEND-1B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 380
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                            SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS
                            CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: BROWDY, Roger L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: HER
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0 CURRENT APPLICATION DATA:
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257 LTVLGLEPPTP 267
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ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
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                                                                FILING DATE:
                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                             CITY: Washington
                                                                                                                                                                                                                                                              STREET:
                  APPLICATION NUMBER:
                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                  STATE:
                                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 419 Seven CITY: Washington STATE: D.C.
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9 Seventh Street, N.W., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (202) 628-5197
202) 737-3528
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UMBER: US 07/854,644
08-SEP-1992
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72.7%;
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Pred. No.
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                                                                                                              Version #1.25
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ATTORNEY/AGENT INFORMATION:

YUN, Allen

REGISTRATION NUMBER:

37,971

HERCEND-1A

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US-08-474-988B-7
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Best Local Similarity 72.7
Matches 8; Conservative
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Patent No. 5874250
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                                               TELEFAX: (202) 737-35
NFORMATION FOR SEQ ID NO:
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                                                                                                                                                          FILING DATE: 04-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION UDATA:
APPLICATION UDATE: US 07/854,644
FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                SEQUENCE CHARACTERISTICS:
                                                                                              REFERENCE/DOCKET NUMBER: HE TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
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                                                                                                                              NAME: YUN, Allen C. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/0 FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
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                                                                              TELEPHONE:
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               ENGTH:
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: 470 amino acids amino acid
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SYSTEM: PC-DOS/MS-DOS
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737-3528
737-7:
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737-3528
70: 7:
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72.7%;
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                                                                                                                              37,971
                                                                                                                HERCEND=1B
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                                                                                                                                                                                                                                                                                                                                                                Version #1
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US-08-416-478A-2
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US-08-394-442B-7
                                                                                                                                        Sequence 2, Application US/08416478A Patent No. 5773578
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Best Local
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                                                                                                                            GENERAL INFORMATION:
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APPLICATION NUMBER: US,
FILING DATE: 24-FEB-199
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                                               TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                              APPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: BROWDY, Roger L. REGISTRATION NUMBER: 25,618
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Triebel;
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                                                New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
AND NEIMARK, P.L.L.C.
                                                                                              Frederic
                                                                                                              Thierry
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Lymphocytes, DNA Sequences Encoding These Proteins And
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    Mismatches

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Pred. No.
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Pred. No.
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Query Match
Best Local Similarity
Best Action 8; Conserve
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                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                          ENERAL INFORMATION:
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                                                                                                                                                                                             ORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 419 Seventh Street, N.W., Suite 400
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/854,644
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/08/474,988B FILING DATE: 07-JUN-1995
                                                                                                                                                                      CITY: Washington STATE: D.C.
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OTHER INFORMATION:
OTHER INFORMATION:
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REGISTRATION NUMBER: 37,971
REFERENCE/DOCKET NUMBER: HE
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                                                                                                                                                                                                                                                            NEW PROTEINS: PRODUCED BY HUMAN LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS AND THEIR PHARMACEUTICAL AND BIOLOGICAL USES
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Pred. No.
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US-08-394-442B-2
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Best Local Similarity
   TELEFAX: (202) 737-35
INFORMATION FOR SEQ ID NO:
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                                               NAME: BROWDY. ROGER L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: HEI
TELECOMMUNICATION INFORMATION:
                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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CORRESPONDENCE ADDRESS
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TITLE OF INVENTION:
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LENGTH: 471 amino acid
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APPLICATION NUMBER:
                                                                                                                               FILING DATE: 24-FEB
CLASSIFICATION: 424
                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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REGISTRATION NUMBER: 37,971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US 07/854,644 FILING DATE: 08-SEP-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 POPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
                                                                                                                                                                                                                                                                               20004
                                                                                                                                                                                                                                                                                                                          Washington
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                                                                                                                                                                                                                                                                                                            D.C.
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                                                                                                                                                                                                                                                                                                                                         E: BROWDY AND NEIMARK, P.L.L.C. 419 Seventh Street, N.W., Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         471 amino acids
                                                                                                                                                                                                                                                                                             U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Triebel, Frederic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                              PatentIn Release #1.0, Version #1.25
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                                 (202) 628-5197
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24-FEB-1995
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2) 620
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MO: 2:
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737-00: 2:
                                                                                                                                                                                                                                                                                                                                                                                                   New Proteins Produced By Human
Lymphocytes, DNA Sequences Encoding
Their Pharmaceutical And Biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65.0%;
72.7%;
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                                                                  HERCEND=2A.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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Matches 8; Conserv
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                                                                                                        TELEX: 248425 EMBON NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                APPLICATION NUMBER: FR 94/05643
FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FR95/00593
FILING DATE: 05-MAY-1995
                                                                                                                                    TELEFORMUNICATION INFORMATION:
TELEPHONE: 703-521-2297
TELEFAX: 703-685-0573
                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 536 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/737,271
                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
OLECULE TYPE:
                                                                                                                                                                                                                       FILING DATE: 05-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: PATCH, Andrew J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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230 LTVLGLEPPTP 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           UMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
LOCATION:
                                         STRANDEDNESS:
                                                                                                                                                                                                   NAME: PATCH, Andrew J. REGISTRATION NUMBER: 32,925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Arlington
                            TOPOLOGY:
                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
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                                                                        476 amino acids
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HUARD, Bertrand
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                            linear
           protein
                                    single
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72.7%;
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Pred. No.
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US-09-058-555-1
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Best Local Similarity
Matches 8; Conserv
              Query Match
Best Local Similarity
 Matches
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APPLICANT: FAURE, Florence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/058,555
                                                                                                                                                                                                                                                   NAME: PATCH, ANDREW J.
REGISTRATION NUMBER: 32,925
REFERENCE/DOCKET NUMBER: BE
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                             FILING DATE: 06-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION UNMERR: PCT/FR95/00593
FILING DATE: 05-MAY-1995
ATTORNBY/AGENT INFORMATION:
NAME: DATE: 05-MAY-1995
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                                                                                                                                                                        SEQUENCE CHARACTERISTICS
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                                                                                                           TOPOLOGY:
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                                                                                                                        STRANDEDNESS:
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CITY: Arlington
STATE: Virginia
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                                                                                                                                                          ENGTH: 476 amino acids
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                                                                                                                                           amino acid
                                                                                              TYPE:
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Conservative
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HUARD, Bertrand
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                                                                                                             linear
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                65.0%;
72.7%;
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72.7%;
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 1; Mismatches
                Score 39; DB Pred. No. 17;
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                             DB 4; Length 476;
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Indels
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Gaps
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235 LTVLGLEPPTP 245

2 LQVLGLQLPTP 12

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Query Match
Best Local Similarity
""" 8; Conserve
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                                                                                                                                                                  Sequence 9, Application US/08474988B Patent No. 5874250
                                                                                                                                                SENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07,
FILING DATE: 08-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                      NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS
                                                                   TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (202) 737-3528

FORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                             257 LTVLGLEPPTP 267
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 37,97:
REFERENCE/FOCKET NUMBER: HITELECOMMUNICATION INFORMATION
TELEPHONE: (202) 628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 04-APR-19
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TITLE OF INVENTION:
         STREET:
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                     ADDRESSEE:
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TOPOLOGY: 11
                                                                                                                                                                                                                                                                             2 LQVLGLQLPTP 12
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OPERATING SYSTEM: PC-DOS/MS-DOS
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419 Seventh Street, N.W., Suite 400
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                                                                                                                  Triebel,
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419 Seventh Street, N.W., Suite 400
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                BROWDY AND NEIMARK, P.L.L.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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                                                                                                                                                                                                                                                                                                                                      65.0%;
72.7%;
                                                             LYMPHOCYTES, DNA SEQUENCES ENCODING THESE PROTEINS THEIR PHARMACEUTICAL AND BIOLOGICAL USES
                                                                                         NEW PROTEINS PRODUCED BY HUMAN
                                                                                                                             Thierry
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Lymphocytes, DNA Sequences Encoding These Proteins And
Their Pharmaceutical And Biological Uses
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                                                                                                                                                                                                                                                                                                                                                   Score 39;
                                                                                                                                                                                                                                                                                                                                       Pred. No.
                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                Length 498;
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Query Match
Best Local Similarity
8; Conserve
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                                                                                                                                                                                                                                                                                                                                                            Sequence 9,
Patent No.
                                                                                                                                                                                                                                                                                                                                                  SENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (202) 628-5197
TELEPAX: (202) 737-3528
NFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                          SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
TITLE OF INVENTION:
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 0 FILING DATE: 08-SEP-1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
FILING DATE: 24
CLASSIFICATION:
                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                         257 LTVLGLEPPTP 267
                             APPLICATION NUMBER: US/08/394,442B
                                                                                                                                                         COUNTRY:
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SOFTWARE: PatentT
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419 Seventh Street, N.W., Suite
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Pred. No. 18;
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ATTORNEY/AGENT INFORMATION:

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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 628-5197
TELEPAX: (202) 737-3528
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 498 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: /protein
                                                                   Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 8; Conserv
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Vatent No. 5955593
                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/706,741B
FILING DATE: 09-SEP-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: HOLLAND, DONALD R.
REGISTRATION NUMBER: 35,197
REFERENCE/DOCKET NUMBER: 965017
TELECOMMUNICATION INFORMATION:
TELEPHONE: (314) 727-5188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SENERAL INFORMATION:
                                                                                                                                                                   STRANDEDNESS:
TOPOLOGY: 11:
MOLECULE TYPE:
                                                                                                                                                                                                                                                                        TELEFAX: (314) 727-60 NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 73 amino acids
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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257 LTVLGLEPPTP 267
1 ELQVLGLQLP 10
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34 ELEVLGRELP 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ALDURESSEE: HOWELL & HAFERKAMP, L.C. STREET: 7733 FORSYTH BLVD., SUITE 1400 CITY: ST. LOUIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: UZIP: 63146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: BROWDY, Roger L. REGISTRATION NUMBER: 2
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ENTION: BH3 INTERACTING DOMAIN DEATH AGONIST
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                                                                 Score 36; DB Pred. No. 7.3; 2; Mismatches
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7.3;
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Search completed: June 4, 2001, 12:14:33 Job time: 226 sec

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Minimum DB
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No.
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12.645 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
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## ALIGNMENTS

Fc gamma (IgG) receptor I (high affinity) form b - human
N;Alternate names: CD64
C;Species: Homo sapiens (man)
C;Date: 03-Apr-1992 #sequence\_revision 03-Apr-1992 #text\_change 21-Jan-2000

C; Accession: A41357; S03019

RESULT A41357

R; Allen, J.M.; Seed, B.
R; Allen, J.M.; Seed, B.
R; Allen, J.M.; Seed, B.
R; Allen, J.M.; Seed, B.
R; Allen, J.M.; Seed, B.
R; Allen, J.M.; Seed, B.
R; Allen, J.M.; Seed, B.
R; Seince 243, 378-381, 1989
A; Title: Isolation and expression of functional high-affinity Fc receptor complementa A; Title: Isolation and expression of functional high-affinity Fc receptor complementa A; Telegraphy Fc receptor complementa A; Title: Isolation and expression: A41357; MUID: 89100284
A; Accession: A41357
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have a	A;Cross-references: GB:X14355; GB:M21090; NID:g31333; PIDN:CAA32536.1; PID:g31334
ing printed,	R; Allen, J.M.; Seed, B.
Ď.	Nucleic Acids Res. 16, 11824, 1988
	A; Title: Nucleotide sequence of three cDNAs for the human high affinity Fc receptor (
	S03018; MUID:89098339
	A; Accession: S03019
	A; Molecule type: mRNA
ription	A; Residues: 1-344 <all2></all2>
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amma (IgG) rec	. A; Note: the authors translated the codon ACT for residue 25 as Ala
amma (IgG) rec	C; Superfamily: Fc gamma receptor I; immunoglobulin homology
amma (IgG) rec	C; Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein
3 protein prec	F;117-170/Domain: immunoglobulin homology <imm></imm>
-enhanced anti	
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thetical prote	.0%; Pred. No. 0.0015;
onine synthase	Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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thetical 16 34	
thetical prote	RESULT 2
somal protein	A39878
somal protein	Fc gamma (IgG) receptor I-A (high affinity) precursor - human
able alcohol d	N;Alternate names: CD64
ue-2-related p	C; Species: Homo sapiens (man)
ue 2 protein -	C;Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change 21-Jul-2000
ue-2 protein -	C; Accession: A39878; I70304; B41357; S03018; I57525
ue-2 protein -	R; van de Winkel, J.G.J.; Ernst, L.K.; Anderson, C.L.; Chiu, I.M.
latory protein	J. Biol. Chem. 266, 13449-13455, 1991
thetical prote	A; Title: Gene organization of the human high affinity receptor for IgG, FcgammaRI (CD
ylylcyclase ty	A; Reference number: A39878; MUID:91302383
thetical prote	A;Accession: A39878
ylylcyclase ty	A; Molecule type: DNA
ylate cyclase	A; Residues: 1-374 < VAN>

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N;Contains: Fc-gamma (IGG) receptor I-B
C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence_revision
C;Accession: I55577; I70303
                             A;Reference number: I55577;
A;Accession: I55577
                                                                                    R; Porges, A.J.; Redecha, P.B.; Doebele, R.;
                                                           A; Title: Novel Fc gamma
   \;Molecule
                                                                                                                                                                           N; Alternate
                                                                                                                                                                                            FC gamma (IgG)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Residues: 1-24,'S',26-337,'T',339-374 <AL2>
A;Cross references: EMBL:X14356; NID:931331;
R;Perez, C.; Wietzerbin, J.; Benech, P.D.
Mol. Cell. Biol. 13, 2182-2192, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Nucleotide sequence of three cDNAs A;Reference number: S03018; MUID:89098339 A;Accession: S03018
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                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gene: GDB:FCGR1A; CD64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Molecule type: DNA
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293-313/Domain: transmembrane #status predicte
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Accession: 157525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Molecule type: mRNA
Residues: 1-24,'S',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title: Isolation and expression of functional high-affinity Reference number: A41357; MUID:89100284 Accession: B41157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Allen, J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Title: Novel Fc gamma
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esidues: 1-24,/S',26-337,'T',339-374 <AL1>
ross-references: GB:X14356; GB:M21091; NID:g31331; PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cross-references: GB:L03418; NID:g184840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ence 243, 378-381,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      292/Domain: extracellular #status predicted <EXT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5/Domain: signal sequence #status predicted
                                                                                                                                                                                                                                                                   277 ELQVLGLQLPTP 288
                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                      1 ELQVLGLQLPTP 12
                                                                                                                                                                                                                                                                                                                                                                                                          152,159,163,195,240/Binding
                                                                              Invest.
         preliminary; translated from GB/EMBL/DDBJ
 type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            translated from GB/EMBL/DDBJ
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12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1-8 <RE2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cis-DNA elements
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                                                                                                                                                                      receptor I-B splice form 1 precursor - nes: CD64
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Res. 16, 11824, 1988
                                                                          90, 2102-2109, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1q21-1q21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90, 2102-2109,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GDB:135911; OMIM:146760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GB:S57204;
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                                            eptor I family MUID:93055454
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, 1992
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Pred. No.
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                                                                                                                  02-Jul-1996 #text_change
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                                                                                      Pan,
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0.0017;
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                                                        products
                                                                                        L.C.;
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                                                                                                                                                                                                                                                                                                                              0;
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                                                                                    Salmon, J.E.;
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Salmon, J.E.; Ki
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                                                     mononuclear cells
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                                                                                   Kimberly, R.P.
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Kimberly, R.P.
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                                 Proc.
A; Reference number:
                                              R; Lau,
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                                                                                                                                B34421
                                                                                                                                             RESULT
                                                                                                                                                                                                                                                                                                                                      C; Keywords:
                                                                                                                                                                                                                                                                                                                                                                                                    A; Gene:
                                                                                                                                                                                                                                                                                                                                                                                                                   C; Genetics:
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S11246
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A; Title: Male-enhanced antigen
                                                               C; Accession: B34421
                                                                           C:Species: Mus musculus (house mouse)
C:Date: 15-Jun-1990 #sequence_revision
                                                                                                         male-enhanced antigen - mouse
                                                                                                                                                                                                                                                                                                                                                   A; Map position: 12p13.3-12pC; Superfamily: human LAG-3
                                                                                                                                                                                                                                                                                                                                                                                 A; Cross-references: GDB:127449;
                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:X51985
A;Note: the author translated the
                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Homo sapiens (man)
C;Date: 18-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 04-Mar-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Accession: S11246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Superfamily: Fc gamma receptor I; immunoglobulin homology C;Keywords: alternative splicing; glycoprotein; immunoglobu F;117-170/Domain: immunoglobulin homology <IMM>
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A;Residues: 1-10,103-153,'A',155-280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GB:L03420;
                              Natl. Acad. Sci. U.S.A. 86,
                                                                                                                                                                                                                      2 LQVLGLQLPTP
                                                                                                                                                                                                                                                                                                                                                                                                    GDB: LAG3
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         precursor
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                                                                                                                                                                                                                                                       Conservative
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s: GB:L03419; NID:g182460; PIDN:AAA35825.1; rm B1
A34421; MUID:90046817
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             kes, R.
86, 8462-8466,
n gene is phylog
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                                                                                                                                                                                                                                                    Score 39; DB
Pred. No. 19;
1; Mismatches
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Pred. No. 6.5;
0; Mismatches
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1.5;
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                                                                         #text_change
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             conserved
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hypothetical protein XF2283 [imported] - Xylella fastidiosa (strain 9a5c) C;Species: Xylella fastidiosa C;Date: 18 Aug-2000 #sequence_revision 20-Aug-2000 #text_change 20-Aug-20
                                                                                                                                   В
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                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Lau,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Homo sapiens (man)
C;Date: 15-Jun-1990 #sequence_revision 15-Jun-1990 #text_change
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A;Residues: 1-174 <KON>
A;Coss-references: EMBL:D17340; NID:g560052; PIDN:BAA04158.1;
A;Note: it is uncertain whether Met-1 or Met-11 is the initiato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Accession: S39790
R;Kondo, M.; Sato, S.; Sutou, S.
Biochim Biophys. Acta 1216, 483-486, 1993
A;Title: Cloning and sequence analysis of cDNA encoding the A;Reference number: S39790; MUID:94092744
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                                                                           RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          male-enhanced antigen - human
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                                                                                                                                                                                                                                                                                                 A;Cross-references: GB:M27937; NID:g187507; PIDN:AAA36208.1; PID:g307182
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                                                                                                                                                                                                                                                                                                                                                                                           ; Reference number: A34421;
                                                                                                                                                                                                                                                                                                                                                                          Accession: A34421
                                                                                                                                                                                                                                                                                                                                                                                           ;Lau, Y.F.C.; Chan, K.; Sparkes, R.
roc. Natl. Acad. Sci. U.S.A. 88, 8462-8466, 1989
roc. Matl. Acad. Sci. U.S.A. 88, 8462-8466, 1989
;Title: Male-enhanced antigen gene is phylogenetically conserved
;Reference number: A34421; MUID:90046817
                                                                                                                                                                                                                                                                                                                                                          ;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Accession: A34421
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                                                                                                                                                                                                                                                                                                                      Molecule type: mRNA
Residues: 1-185 <LAU>
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Best Local
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Best Local
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Best Local
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100 IQALGLHLPDP 110
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                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity 63.0
7; Conservative
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#sequence_revision 20-Aug-2000 #text_change 20-Aug-2000
                                                                                                                                                                                                                       63.3%;
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Pred. No.
                                                                                                                                                                                                                       Score 38; DB 2; Pred. No. 9.6;
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Pred. No. 9;
                                                                                                                                                                                                          Mismatches
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Query Match
Best Local Similarity
--hes 6; Conserv
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A;Note: for a complete list of authors see reference number A59328 below
                                                                                                                                        C;Superfamily: threonine synthase
C;Keywords: carbon-oxygen lyase; phosphoprotein;
F;105/Binding site: pyridoxal phosphate (Lys) (co
                                                                                                                                                                                                                                                                                                                                                   A;Authors: Gnehm, C.L.; McDonald, L.A.; Small, K.V.; Fraser, C.M.; Smith, H.O.; Vente A;Title: Whole-genome random sequencing and assembly of Haemophilus influenzae Rd. A;Reference number: A64000; MUID:95350630
A;Accession: H64047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Experimental source: strain 9a5c
R;Simpson, A.J.G.; Reinach, F.C.; Arruda
Briones, M.R.S.; Bueno, M.R.P.; Camargo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R; anonymous, The Xy
Nature 406, 151-157
                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-425 <TIGR>
                                                                                                                                                                                                                                                                                                                        A;Status: .nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Science 269, 496-512, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Fleischmann, R.D.; Adams, M.D.; White, O.; Clayton, R.A.; Kirkness, E.F.; Ker; Gocayne, J.D.; Scott, J.; Shirley, R.; Liu, L.I.; Glodek, A.; Kelley, J.M.; V. D.M.; Brandon, R.C.; Fine, L.D.; Fritchman, J.L.; Fuhrmann, J.L.; Geoghagen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Date:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                threonine synthase (EC 4.2.99.2) - Haemophilus influenzae (strain Rd C;Species: Haemophilus influenzae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, M.C.; de Oliveira, M.C.; de Sa, R.; Rodrigues, V.; Rosa, A.J. de M.; de Rosa Jr., V.E.; de Sa, R. A.M.; A.K.; da Silva, F.R.; da Silva, A.M.; M.; Tsuhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Al
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to GenBank, June 2000

A; Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.;

A; Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.P.; Krieger, J.E.; Kuramae, E.E.;
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                                                                                                                                                                                                                                A;Cross-references: GB:U32694; GB:L42023; NID:g1573035; PIDN:AAC21765.1; PID:g1573038
   Matches
                                 Query Match
Best Local
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                              Local
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Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         H64047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Docena, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J
                           63.3%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R. amargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer
                           Score 38; DB Pred. No. 24;
Mismatches
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                                                                                                                                  oprotein; pyridoxal phosphate; threonine biosy (Lys) (covalent) #status predicted
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17;
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ilva, A.M.; Silva Jr., W.A.; da
Verjovski-Almeida, S.; Vettore,
   1;
                                                         Length 425;
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   Indels
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Ferreira, A.J.S.
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   Gaps
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ne, E.E.; La
V.; Martins
Miyaki, C.
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M.; Weidman
Jen, N.S.M.
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386 RILGIQLPLP 395

3 QVLGLQLPTP 12

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M: Shen, M.; Vamathevan, J.J.; Lam, S.; Smith, H.O.; Venter, J.C.; Fraser, Science 286, 1571-1577, 1999
                                              A; Title: Genome sequence of A; Reference number: A75250;
                                                                                                                                                                transcription regulator, IclR family C;Species: Deinococcus radiodurans C;Date: 03-Dec_1999 #sequence_revision
                                   A; Accession:
                                                                                                                                 R; White, O.; Eisen,
            \;Status: preliminary
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Best Local Similarity
Thes 7; Conserve
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Cience 282, 754-759, 1998
Cience 282, 754-759, 1998
Cience 282, 754-759, 1998
Cience 282, 754-759, 1998
Cience 282, 754-759, MUID:99000809

A.Accession: E71475
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                                                                                                                                                     Accession: C75611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               submitted to the EMBL Data Library,
                                                                                                                                                                                                                                                                                                                                                                                                                             Superfamily: phosphoglycerate mutase; phosphoglycerate mutase homology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         robable phosphoglycerate mutase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cross-references: GB:AE001343; GB:AE001273; NID:g3329176; PIDN:AAC68317.1; PID:g33291:
Experimental source: serotype D, strain UW-3/Cx
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Date: 13-Sep-1998 #sequence_revision 13-Sep-1998 #text_change 16-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           species:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cross-references: EMBL:AF043
Experimental source: strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: DNA Residues: 1-134 <TIN>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Status: preliminary; translated from GB/EMBL/DDBJ
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                                                                                                                                                                                                                                                                                  195 EEQVLSLELPT 205
                                                                                                                                                                                                                                                                                                                                                         Local Similarity
les 8; Conserv
                                                                                                                                                                                                                                                                                                                     1 ELQVLGLQLPT 11
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                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                J.A.; Heidelberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL:AF043698; PIDN:AAB97562.1; GSPDB:GN00019; CESP:C54G6.4
                                                                                                                                                                                                                                                                                                                                                                    61.7%;
72.7%;
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70.0%;
                                              MUID:20036896
                                                          the radioresistant bacterium
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Pred. NO. 18;
1; Mismatches
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Pred. No. 10;
1; Mismatches
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                                                                                                                              J.F.;
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                                                                                                                                                              03-Dec-1999
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                                                                                                            , Hickey, I
McDonald,
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                                                                                                       E.K.;
1, L.;
                                                                                                                                                          #text_change 17-Mar-2000
                                                                                                                                                                                          radiodurans (strain R1)
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                                                                                                                                                                                                                                                                                                                                                                                Length 226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                        Deinococcus
                                                                                                       ; Peterson, J.D.; Dodson, Utterback, T.; Zalewski,
                                                                                                                                                                                                                                                                                                                                                  0;
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                                                                                                       R.J.;
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Query Match
Best Local Similarity
                                                                                                       ; Gene :
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A; Map position:
                                                                                       A; Experimental source: clone T08G11
                                                                                                                                                                                        submitted to the RMBL Data Library, September 1996
A;Reference number: Z19925
A;Accession: T24692
                                                                                                                                                                                                                                                               C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_rev
C;Accession: T24692
                                                                                                    A; Cross-references: EMBL: Z80220; PIDN: CAB02304.1;
                                                                                                                          A; Molecule type: DNA
A; Residues: 1-3212 <WIL>
                                                                                                                                                           A; Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                   R;Dobson,
                                                                                                                                                                                                                                                                                                                          hypothetical protein T08G11.1 -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Introns: 154/1; 326/3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    submitted to the EMBL Data Library, A;Description: The sequence of C. e A;Reference number: Z21420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Map position: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: EMBL:AF106584; PIDN:AAC78220.1; A;Experimental source: strain Bristol N2; clone F54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-424 <JON>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Gene: CESP:F54A5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Genetics:
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Best Local Similarity
Trhes 7; Conserv
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C;Superfamily: acetate operon repressor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-253 <WHI>
                                                                           Genetics:
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Best Local Similarity 70.0%;
Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species:
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                                                 CESP:T08G11.1
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            149/3;
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  182/3; 527/2; 766/1; 1070/3; 1769/3; 1834/3; 2070/2;
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Pred. No.
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Y, November 1998
elegans cosmid F54A5
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Pred. No.
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F54A5
                                                                                                    GSPDB:GN00019; CESP:T08G11
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61.7%;

Score Pred.

37;

DB ·2; 3.5e+02;

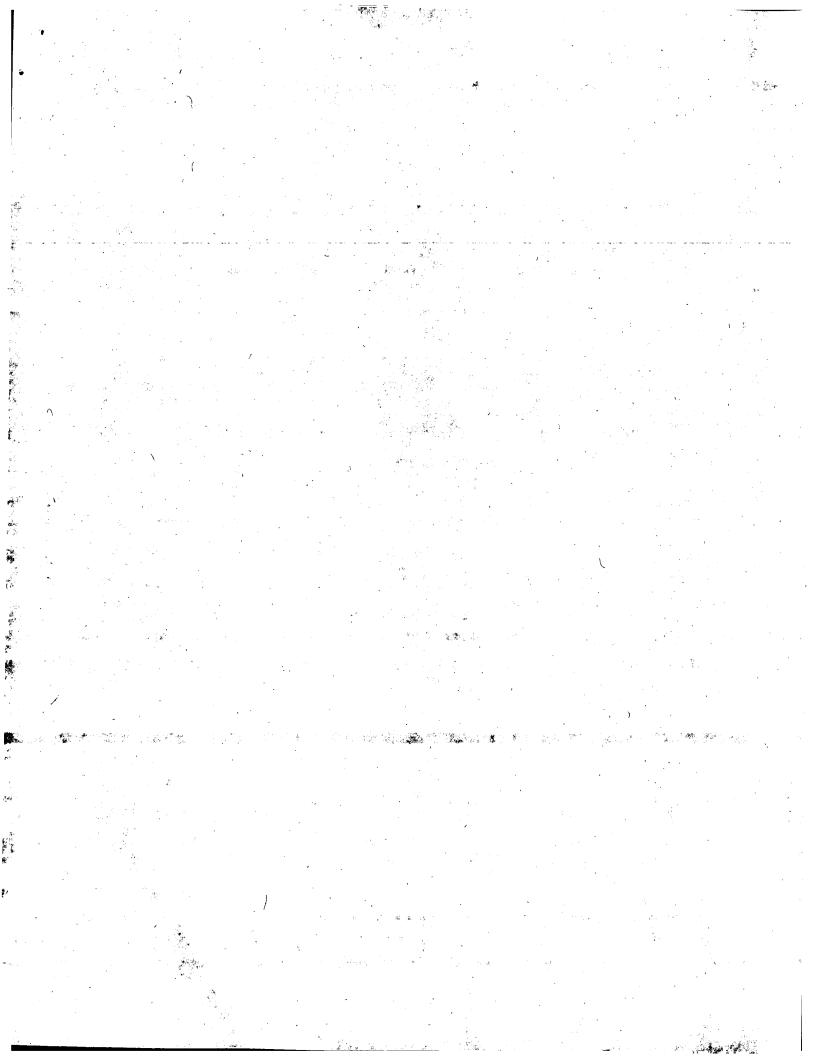
Length 3212

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Search completed: June 4, 2001, 12:15:46 Job time: 279 sec
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                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: DNA
A;Residues: 1-144 <HES>
A;Residues: 1-144 <HES>
A;Cross-references: EMBL:X84724; NID:g780165; PIDN:CAA59211.1; PID:g780172
A;Cross-references: EMBL:X84724; NID:g780165; PIDN:CAA59211.1; PID:g780172
R;Hess, M; Cuzange, A; Ruigrok, R.W.H.; Chroboczek, J;; Jacrot, B.
J; Mol. Biol. 252, 379-385, 1995
A;Title: The avian adenovirus penton: two fibres and one base.
A;Reference number: S59067; MUID:96025073
A;Contents: annotation
C;Superfamily: fowl adenovirus 1 hypothetical 16.34K protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nypothetical 16.34K protein - fowl adenovirus 1

);Species: Aviadenovirus gall (fowl adenovirus 1, CELO)

;Date: 08-Jul-1995 #sequence_revision 14-Feb-1997 #text_change 22-Oct-1999

;Accession: S54130
                                                                                                                                                                                                                                                   Query Match 60.0%;
Best Local Similarity 85.7%;
Matches 6; Conservative
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772 LMKLGLSIPTP 782
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39 GLQMPTP 45
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1; Mismatches 0; Indels
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4.5 Compugen Ltd.

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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FCG1_MOUSE
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PRTZ_HORVU
        KG3A_HUMAN
PT1_STAAU
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RL1X_HUMAN
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THRC_HAEIN
P2X2_CAVPO
RL20_YEAST
BID_MOUSE
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	salmonella	salmonella	escherichia		escherichia	_	chlamydia t	chlamydia m	methanococc	saccharomyc	halocynthia	schizosacch	,

## ALIGNMENTS

EMBL; X14356; CAA32537.1; EMBL; X14355; CAA32536.1;

PIR;

PIR; B41357; MIM; 146760;

1ALT:

Pro; IPR003006; PF00047; ig; 3

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; S03019; S03019; ; A41357; A41357; ; B41357; B41357. S03018; S03018

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MEDLINE=90237736; Pubmed=1692078;
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MEDLINE=90237736; Pubmed=1692078;
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Viegas-Pequignot E., F
"LAG-3, a novel lympho"
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_HUMAN
LAG3_HUMAN
P18627;
                                                                    Hercend
                                                                                           MEDLINE=92364535;
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                                                        "Characterization
                                                                                                                               Submitted
                                                                                                                                             Triebel
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Eukaryota; Metazoa; C
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Pred. No: 0.0
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Catarrhini; Hominidae; Homo.
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TISSUE-Small intestine; Winteroe A.K., Fredholm M., Davies W.; Winteroe A.K., Fredholm M., Davies W.; Submitted (OCT-1996) to the EMBL/GenBank/DDBJ databases -1- FUNCTION: MAY PLAY AN IMPORTANT ROLE IN SPERMATOGENI TESTIS DEVELOPMENT.
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         Sus scrofa (Pig).
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                            MALE-ENHANCED
MEA1 OR MEA.
NCBI_TaxID=9823;
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EMBL; A21353; CAA01547.1;
PIR; S11246; S11246.
MIM; 153337; -
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ANTIGENS.
- SUBCELLULAR LOCATION: 7
- TISSUE SPECIFICITY: ON
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8; Conservative
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MEAL OR MEA.
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                                                                                                                            Spermatogenesis; Developmental protein. SEQUENCE 172 AA; 18544 MW; 9342AFAD2C96FF25 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                      "Male-enhanced antigen gene is phylogenetically conserved and expressed at late stages of spermatogenesis.";
Proc. Natl. Acad. Sci. U.S.A. 86:8462-8466(1989).
-I- FUNCTION: MAY PLAY AN IMPORTANT ROLE IN SPERMATOGENESIS AND/OR
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NCBI_TaxID=9606;
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TISSUE SPECIFICITY: HIGHLY EXPRESSED IN TESTIS.
CAUTION: WAS ORIGINALLY THOUGHT TO BE THE H-Y ANTIGEN.
CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-11 IS THE INITIATOR.
                                                                                                                                                                                                                                                         ween the Swiss Institute of Bioinformatics and the EMBL outstation
European Bioinformatics Institute. There are no restrictions on it
by non-profit institutions as long as its content is in no wa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    35,
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                                                                                                                                                                                                                                                                                                                                                                                                                                              K., Sparkes R.S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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                                                      Score 38; DB Pred. No. 3.7; 1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1F52DC82FDAA95B6 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   update)
                                                                     DB 1;
3.7;
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                                                                                   Length 172;
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RESULT 6
MEA1_MOUSE
ID MEA1_M
AC Q64327
DT 01-NOV
DT 01-NOV
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Best Local
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                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by an
entities requires a license agreement (See http://www.isb-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Cloning and sequence analysis of cDNA encoding derived male-enhanced antigen (Mea)."; Biochim. Biophys. Acta 1216:483-486(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1997 (Rel. 35, Last sequence update) 01-NOV-1997 (Rel. 35, Last annotation update) MALE-ENHANCED ANTIGEN-1 (MEA-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEA1_BOVIN Q29407;
                                                                                                                                                                                                                                                                                             EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       during spermatogenesis.";
DNA Seq. 6:75-85(1996).
-!- FUNCTION: MAY PLAY AN IMPORTANT ROLE IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Che
Mammalia; Eutheria; Ce
Bovidae; Bovinae; Bos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1997
01-NOV-1997
                                                                                                                                                                                                                                                                Spermatogenesis; Description of the Sequence 174 AA;
                                                                                                                                                                                                                                                                                                            EMBL;
                                                                                                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIINE=97063437; PubMed=8907304;
Kondo M., Terouchi S., Tsukasa N., Sato.S., Ishida N., Sutou S.;
"Genomic sequence analysis of the bovine male-enhanced antigen-1
(Mea-1) and differential localization of its transcripts and pro-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9913;
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                                                                                                                                             2 LQVLGLQLPTP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          CAUTION:
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TISSUE SPECIFICITY: HIGHLY EXPRESSED IN TESTIS. TRANSCRIPTS CAN BE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          THE PROTEIN ITSELF IS ONLY DETECTED IN SPERMATIDS. NO EXPRESSION IN LETYDIG CELLS, SPERMATOGONIA, OR SPERM. VERY WEAK EXPRESSION IN THE HEART, KIDNEY, SPLEEN, THYMUS, AND OVARY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DEVELOPMENTAL STAGE: EXPRESSED IN THE TO HATCHED BLASTOCYTS.
                                                                                                                                                                                                                                                                                          D17340; BAA04158.1; D30811; BAA06488.1;
                                                                                                                                                                                       Similarity 7; Conserv
                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                         IT IS UNCERTAIN WHETHER MET-1 OR MET-11 IS THE INITIATOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
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35, Last sequences
35, Last annotations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                      63.3%;
                                                                                                                                                                                                      Score 38; I
Pred. No. 3.
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Bd
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01-NOV-1997 01-NOV-1997 01-NOV-1997

(Rel. (Rel. (Rel.

35, Created)35, Last sequence update)35, Last annotation update)

MEA1\_MOUSE Q64327;

STANDARD;

174

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Best Local :
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01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NOV-1995 (Rel. 01-NO
                                                                                           Haemophilus influenzae.
Bacteria; Proteobacteria;
Haemophilus.
                                                                                                                                                                               THRC OR HI0087
                                                                                                                                                                                                                                                                                                                                                                    HAEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Spermatogenesis; Developmental SEQUENCE 174 AA; 18584 MW;
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EMBL; L10401; AAA39520.1; -.
EMBL; D17341; BAA04159.1; -.
MGD; MGI:96957; Meal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See or send an email to license@isb-sib.ch).
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Biochim. Biophys. Acta 1216:483-486(1993).
-i-FUNCTION: MAY PLAY AN IMPORTANT ROLE IN SPERMATOGENESIS AND/OR TESTIS DEVELOPMENT.
-i-TISSUE SPECIFICITY: HIGHLY EXPRESSED IN TESTIS. TRANSCRIPTS CAP.
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MEDLINE=90046817; PubMed=2813404;
Lau Y.-F.C., Chan K., Sparkes R.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Male-enhanced antigen gene is phylogenetically conserved expressed at late stages of spermatogenesis."; Proc. Natl. Acad. Sci. U.S.A. 86:8462-8466(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
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MEA1 OR MEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 LQVLGLQLPTP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FOUND IN PRIMARY AND SECONDARY SPERMATOCYTES, AND SPERMATIDS, BUT THE PROTEIN ITSELF IS ONLY DETECTED IN SPERMATIS. NO EXPRESSION IN LEYDIG CELLS, SPERMATOGONIA, OR SPERM. VERY WEAK EXPRESSION IN THE HEART, KIDNEY, SPLEEN, THYMUS, AND OVARY.

DEVELOPMENTAL STAGE: EXPRESSED AS EARLY AS DAY 6 POST-PARTUM, WITH HIGHER EXPRESSION IN THE ADULT TESTIS.

CAUTION: IT IS UNCERTAIN WHETHER MET-1 OR MET-11 IS THE INITIATOR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IQALGLHLPDP 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SWISS-PROT entry is copyright. It is produced through a en the Swiss Institute of Bioinformatics and the EMBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity 7; Conserv
                                                                                                                                                                                                        SYNTHASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sato
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                                                                                                                                                                                                                              32, Created)
32, Last sequences
32, Last annotations
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Rodentia;
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                                                                                                                       gamma
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C3B16361635B176F CRC64;
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Sciurognathi; Muridae; Murinae; Mus
                                                                                                                  subdivision;
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on update)
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Best Local :
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                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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SMOOTH MUSCLE.
SUBUNIT: HOMO-
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P2X2_CAVPO

970397; 070398;

15-DEC-1998 (Rel. 37, C

15-DEC-1998 (Rel. 37, L

30_MAY-2000 (Rel. 39, L
                                                                                                                   STRAIN=NIH 2; TISSUE=Organ of corti;
MEDLINE=98346338; PubMed=9682808;
Parker M.S., Larroque M.L., Campbell
                                                                                                                                                                                                                                       Cavia porcellus (Guinea pig).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS00165; DEHYDRATASE_SER_THR; 1.
Threonine biosynthesis; Lyase; Pyridoxal phosphate.
BINDING 105 105 PYRIDOXAL PHOSPHATE (BY S. SEQUENCE 425 AA; 46682 MW; 2CC222DD201F09E0 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR000634; -.
InterPro; IPR001926; -.
Pfam; PF00291; S_T_dehydratase; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or send an email to license@isb-sib.ch)
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-!- PATHWAY: THREONINE BIOSYNTHESIS
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  FUNCTION: BINDING MEDIATES SYNAPTIC
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                                            Res. 121:62-70(1998)
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A.R., Bu
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                                                                                    of the P2X2
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OF THIS LIGAND GATED TRANSMISSION BETWEEN
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                                                                               receptor
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T., Hedblom E., Cotton M.D.,
.T., Saudek D.M., Brandon R.C.
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  ION CHANNEL NEURONS AND
                                                                                                                   Bobbin R.P.,
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Best Local
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SEQUENCE FROM N.A. (RESEQUENCE FROM N.A.
                                                                                                               Gentles S. Submitted
                                                                                                                                                                                                                                                         (RPL20A OR RPL18A2 OR YMR242C OR YM9408.04C) AND (RPL20B OR RPL18A1 OR RPL18B OR YOR312C OR O6116). Saccharomyces cerevisiae (Baker's yeast).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by an
                               Pearson B.M., Hernando Y., Wolf S.S.,
                                                                                                                                                                                                                                                                                                                                     01-FEB-1996 (Rel.
15-DEC-1998 (Rel.
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Tonic channel; Transmembrane; Ion transport; Receptor; Glycoprotein;
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                     Schweizer M.
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InterPro; IPR003045; -
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(AUG-1995) to the EMBL/GenBank/DDBJ databases
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or send a
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P70444;
                                                                                                                                                                                                                McDonnell J.M., Fushman D., Milliman C.L., Korsme "Solution structure of the proapoptotic molecule basis for apoptotic agonists and antagonists."; Cell 96:625-634(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; Z48756; CAA88652.1; ALT_SEQ.
EMBL; X90565; CAA62167.1; -.
EMBL; Z75220; CAA99632.1; -.
SGD; S0004855; RPL20A.
SGD; S0005839; RPL20B.
                                                                                                                                                                                                                                                                                                                Wang K., Yin X.-M., Chao D.T., Milliman C.L., "BID: a novel BH3 domain-only death agonist."; Genes Dev. 10:2859-2869(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
30-NAY-2000 (Rel. 39, Last annotation updat
BH3 INTERACTING DOMAIN DEATH AGONIST (BID).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ribosomal protein; Multigene family
SEQUENCE 174 AA; 20713 MW; 52BA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produbetween the Swiss Institute of Bioinformatics the European Bioinformatics Institute. There \varepsilon
                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                            -!- SUBUNIT: FORMS HETERODIMERS EITHER W
BAX OF THE ANTI-APOPTOTIC PROTEIN BC
-!- SUBCELULAR LOCATION: PREDOMINANTLY
-!- DOMAIN: INTACT BH3 DOMAIN IS REQUIRE
                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR002670; -.
Pfam; PF01775; Ribosomal_L18ae;
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                                                                                                                                                                                     EFFECT OF BCL-2.
                                                                                                                                                                                                                                                                          MEDLINE=99189748; PubMed=10089878;
                                                                                                                                                                                                                                                                                         STRUCTURE BY NMR.
                                                                                                                                                                                                                                                                                                                                                               MEDLINE=97078762; PubMed=8918887;
                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- MISCELLANEOUS: THERE ARE TWO GENES FOR L20
-!- SIMILARITY: BELONGS TO THE L18AE FAMILY OF
                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A., AND MUTAGENESIS OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10
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                                                                                  APOPTOTIC MEMBERS OF THE BCL-2 FAMILY.
SIMILARITY: CONTAINS A BCL-2 HOMOLOGY DOMAIN 3 (BH3)
                                                                                                             DOMAIN: INTACT BH3 DOMAIN IS REQUIRED BY BIK, BID, BAI THEIR KILLING ACTIVITY AND FOR THEIR INTERACTION WITH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           s requires a license agreement (
an email to license@isb-sib.ch)
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               52BA8648F9531208 CRC64;
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No. 9.1;
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(BID)
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                                                                                                                                                                                                                                                                                                                                                                                            BH3 DOMAIN
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                                                                                                                            CYTOPLASMIC.
FD BY BIK, BID,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       as
                                                                                                                                                                                                                                             Korsmeyer S.J., Cowburn lecule BID: a structural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                       THE PRO-APOPTOTIC PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Usage by
                                                                                                                                                                                                                                                                                                                                                 Korsmeyer S.J.;
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RIBOSOMAL PROTEINS.
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RESULT 12
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Matches 7
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PROSITE; PS00518; ZINC_FINGER_C3HC4; UNKNOWN_1.
                                                                                                                                                                                                                   use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                   Hypothetical protein.
SEQUENCE 292 AA; 2
                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a continuous the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Y161_HUMAN
P50876;
-01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _HUMAN
                                                                                                                                                                                                 EMBL; D79983; BAA11478.1;
                                        121
                                                                                                                                                                                                                                                                                                        "Prediction of the coding sequences of unidentified human genes."
The coding sequences of unidentified human genes.
The coding sequences of 40 new genes (KIAA0161-KIAA0200) deduced analysis of cDNA clones from human cell line KG-1.";
DNA Res. 3:17-24(1996).
                                                                                                                                                                                                                                                                                                                                                         TISSUE=Bone marrow;
MEDLINE=96281124; PubMed=8724849;
                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL
                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens (Human)
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DOMAIN 86 100
SEQUENCE 195 AA; 21950 MW;
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InterPro; IPR000712;
PROSITE; PS01259; BH3; 1.
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                                                           1 ELQVLGLQLPTP 12
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(Rel.
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. 34, Last sequence. 40, Last anno
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                                                                                                                                 32859 MW;
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Primates;
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50
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70.0%;
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Pred. No.
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                                                                                                Length 292;
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                    CONFLICT
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P12959;
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                                                                                                                                    T00668;
        453 AA;
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PROSITE; PS00036; BZIP_BASIC; Nuclear protein; DNA-binding;
                                                                                                                                                                                                    InterPro; IPR001871; -
Pfam; PF00170; bZIP; 1
                                                                                                                                                                                                                                                                                     EMBL; X15544; CAA33550.1;
EMBL; X16618; CAA34614.1;
PIR; S06022; S06022.
HSSP; P03069; 2DGC.
                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a license agreement
                                                                                                                                                                                                                                                                                                                                                                                                                                use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Salamini F., Thompson R.D.;
"The maize regulatory locus Opaque-2 encodes a DNA-binding which activates the transcription of the b-32 gene.";
EMBO J. 10:617-624(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "The O2 gene which regulates zein encodes a protein with structural activators.";
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MEDLINE=90016825; PubMed=2798113;
Maddaloni M., di Fonzo N., Hartings H., La
Salamini F., Thompson R.D., Motto M.;
"The sequence of the zein regulatory gene ,
Nucleic Acids Res. 17:7532-7532(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hartings H., Maddaloni M., Salamini F., Thompson R.D.
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STRAIN-AC 1503 GM 1407;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=91160516; PubMed=2001677; Lohmer S., Maddaloni M., Motto M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBO J. 8:2795-2801(1989).
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Eukaryota; Viridiplantae; 1
Magnoliophyta; Liliopsida;
Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JAN-1990 (Rel. 13, Create 01-JAN-1990 (Rel. 13, Last s 01-FEB-1996 (Rel. 33, Last a 02-AQUE-2 REGULATORY PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: TO OTHER BZIP PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: INVOLVED IN THE REGULATION OF THE ENDOSPERM-SPECIFIC PRODUCTION OF ALBUMIN B-32 AND OTHER ZEIN PROTEINS. IT IS A TRANS-ACTING TRANSCRIPTIONAL ACTIVATOR THAT BINDS TO THE CONSENSUS SEQUENCE 5 '-GATGAYRTGR-3'.
SUBCELLULAR LOCATION: NUCLEAR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SWISS-PROT entry is copyright. It is produced through
een the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thompson R.D.
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                   LEUCINE-ZIPPER.
POTENTIAL.
E -> EPEPEPE (IN
D -> A (IN REF.
K -> KR (IN REF.
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                                                                                                                       BASIC MOTIF
                                                                                                                                                         Transcription regulation;
                                    -> EPEPEPE (IN
513A8AB8D5ABD999
                     KR (IN REF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         deposition homologies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   di Fonzo N., Motto
                                                                                                                                                                                                                                                                                                                                                                                                http://www.isb-sib.ch/announce/
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                                                           REF.
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transcriptional
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                                                                                                                                                           Activator;
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Query Match Best Local

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Best Local Similarity
Matches 7; Conserv
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                                   Q04400;
01-OCT-1993 (Rel. 2
01-NOV-1997 (Rel. 3
30-MAY-2000 (Rel. 3
                                                                                                  RAT
                                                                                                                                                                                                                                                        ZN_FING
ZN_FING
SEQUENCE
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Q19345;
                                                                                      CYA5_RAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAEEL
                           ADENYLATE CYCLASE
                                                                                                                                                                                                                                                                                            DNA_BIND
                                                                                                                                                                                                                                                                                                                                         Pfam; PF00104; hormone_rec; 1
Pfam; PF00105; zf-C4; 1.
PRINTS; PR00047; STROIDFINGER
                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR000536; -. InterPro; IPR001628; -.
                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; Z54270; CAA91028.1; WormPep; F11C1.6; CE03191.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     entitles requires a license agreement (
or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the European Bioinformatics Institute.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Palmer S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN-BRISTOL N2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Rhabditidae; Pelode;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Caenorhabditis elegans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NHR-25 OR F11C1.6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
NUCLEAR HORMONE RECEPTOR FAMILY MEMBER NHR-25
                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00031;
                                                                                                                                                                                                                                                                                                                                                                                                      HSSP; P19793; 2NLL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=6239;
                                                                                                                                                 551
                                                                                                                                                                                                                                                                                                                   Receptor; Transcription
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                                                                                                                                                                                                                                                                                                         inc-finger.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1- SIMILARITY: BELONGS TO THE NUCLEAR HORMONE RECEPTORS FAMILY.
                                                                                                                                                                     3 QVLGLQLPTP 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ELQVLGLQLPT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mitted (SEP-1995) to the EMBL/GenBank/DDBJ databases FUNCTION: ORPHAN NUCLEAR RECEPTOR. SUBCELLULAR LOCATION: NUCLEAR (POTENTIAL).
                                                                                                                                              QVPAIQLPTP 560
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5; Conserv
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18
54
572 AA;
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                                                                                      STANDARD;
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.27, Created)
35, Last sequence update)
39, Last annotation update)
2, TYPE V (EC 4.6.1.1) (ATP PYROPHOSPHATE-LYASE)
                                                                                                                                                                                                                                                                                                                   NUCLEAR_RECEPTOR; iption regulation;
                                                                                                                                                                                                                                                        83
38
78
64772 MW;
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                                                                                                                                                                                                         70.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 36; DB Pred. No. 25; 5; Mismatches
                                                                                                                                                                                                         Score 36;
Pred. No.
                                                                                                                                                                                                                                                                    NUCLEAR RECEPTOR-TYPE C4-TYPE. C4-TYPE.
                                                                                      PRT;
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                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                           1.
DNA-binding; Nuclear
                                                                                     1262 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      572 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB
25;
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32;
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                                                                                                                                                                                                                     1;
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                                                                                                                                                                                                                    Length 572;
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Query Match Best Local S Matches 7

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Score 36; DB Pred. No. 75; 2; Mismatches

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CARBOHYD
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                               DOMAIN
                                                                                                                                                                                                                           TRANSMEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00211;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; M96159; AAB39764.1; HSSP; P19754; LAWK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between 'the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Premont R.T.; Submitted (DEC-1996) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-SPRAGUE-DAWLEY; TISSUE-Liver, Kidney, and Heart;
                                                                                                                  DOMAIN
                                                                                                                                  TRANSMEM
                                                                                                                                                     TRANSMEM
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                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -1- CATALYTIC ACTIVITY: ATP = 3',5'-CYCLIC AMP + PYROPHOSPHATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Proc. Natl. Acad.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Two members of a widely expressed adenylyl cyclases.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Premont R.T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ADENYLYL CYCLASE).
ADCY5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -!- FUNCTION: THIS A MEMBRANE-BOUND,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=93028552; PubMed=1409703;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE OF 164-1262 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                 yase;
                                                                                                                                                                                                                                                                                                                                                                           RANSMEM
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SUBCELLULAR LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOMAIN: COMPOSED OF TWO HOMOLOGOUS DOMAINS.
SIMILARITY: BELONGS TO ADENYLYL CYCLASE CLASS-4/GUANYLYL CYCLASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FAMILY. ~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. CONTAINS DOMAINS WITH 6 POTENTIAL TRANSMEMBRANE REGIONS EACH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-profit
                                                                                                                1006
                                                                                                                                                                                                                                                                                                                                                                                                             synthesis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   guanylate_cyc;
                                                                                                                                                                                                                                                                                                                                                                                                                               GUANYLATE_CYCLASES_1;
GUANYLATE_CYCLASES_2;
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1262
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     139178
                                                                                                                                                                                                                                                                                                                                                                                                               Transmembrane;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    & GAMMA SUBUNIT COMPLEX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U.S.A. 89:9809-9813(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ma H.-W.,
     Ψ¥.
N-LINKED (GLCNAC. . .)
N-LINKED (GLCNAC. . .)
N-LINKED (GLCNAC. . .)
NW; 8B147E201C5DF601 CI
                                                                                                              CYTOPLASMIC (POTENTIAL).
                                                                                                                                  POTENTIAL
                                                                                                                                                                                                                                                                                                                                                                                            membrane; Glycoprotein; Duplication CYTOPLASMIC (POTENTIAL).
                                                                              POLY-ALA.
                                                                                             POLY-GLN.
                                                                                                                                                     POTENTIAL.
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                                                                                                                                                                                                                                               POTENTIAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CALMODULIN-INSENSITIVE ADENYLYL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (See http://www.isb-sib.ch/announce,
                                                                                                                                                                                                                                                                                                                                                                                                                                NN
                                                                                                                                                                                                                                                              (POTENTIAL)
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hormone-stimulated
   .) (POTENTIAL) CRC64;
                                        (POTENTIAL)
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                     DOMAIN
                                                                                                                                                           NIAMOC
                                                                                                                                                                                                                                                                            Pfam; PF00211; guanylate_cyc; 2.
PROSITE; PS00452; GUANYLATE_CYCLASES_1; 2.
PROSITE; PS50125; GUANYLATE_CYCLASES_2; 2.
                                                                                                                                                                                                                                                                                                             InterPro; IPR001054; Pfam; PF00211; guany
                                                                                                                                                                                                                                                                                                                                          EMBL; Z29371; CAA82562.1; HSSP; P19754; IAWK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE-Myocardium;
MEDLINE-94139935; PubMed=8307190;
Wallach J., Droste M., Kluxen F.-W., Pfeuffer T.,
"Molecular cloning and expression of a novel type
from rabbit myocardium.";
FEBS Lett. 338:257-263(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P40144;
01-FEB-1995 /
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -1- FUNCTION: THIS A MEMBRANE-BOUND, CA(2+)-INHIBITABLE ADENYLYL CYCLASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A., AND PARTIAL SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-FEB-1995 (Rel. 31, Last sequence update)
30 MAY-2000 (Rel. 39, Last annotation update)
ADENYLATE CYCLASE, TYPE V (EC 4.6.1.1) (ATP PYROPHOSPHATE-LYASE)
CADENYLYL CYCLASE).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CYA5_RABIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PTM: THE N-TERMINUS IS BLOCKED.
SIMILARITY: BELONGS TO ADENYLYL CYCLASE CLASS-4/GUANYLYL CYCLASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CONCENTRATION RANGE. SUBCELULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. TISSUE SPECIFICITY: MYCCARDIAL TISSUE. DOMAIN: COMPOSED OF TWO HOMOLOGOUS DOMAINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CATALYTIC ACTIVITY: ATP = 3',5'-CYCLIC AMP + PYROPHOSPHATE. ENZYME REGULATION: INHIBITION BY CA(2+) IN THE SUBMICROMOLAR
                                                                                                                                                                                                                                                                       synthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
816
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82
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                                                                                                                                                                                                                                                             Transmembrane; Glycoprotein; Duplication
                                                                                                                                                                               POTENTIAL.
POTENTIAL.
                                                                                                                          POTENTIAL.
                                      POTENTIAL.
CYTOPLASMIC
                                                                                EXTRACELLULAR POTENTIAL.
                                                                                                                                                   POTENTIAL.
CYTOPLASMIC
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                                                                    POTENTIAL.
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novel type V adenylyl cyclase
                                     (POTENTIAL).
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FT CARBOHYD 873 873 N-LINKED (GLCNAC...) (POTENTIAL).
FT CARBOHYD 890 890 N-LINKED (GLCNAC...) (POTENTIAL).
FT CARBOHYD 975 975 N-LINKED (GLCNAC...) (POTENTIAL).
SQ SEQUENCE 1264 AA; 139623 MW; 1787EB42A0C2FDF6 CRC64;

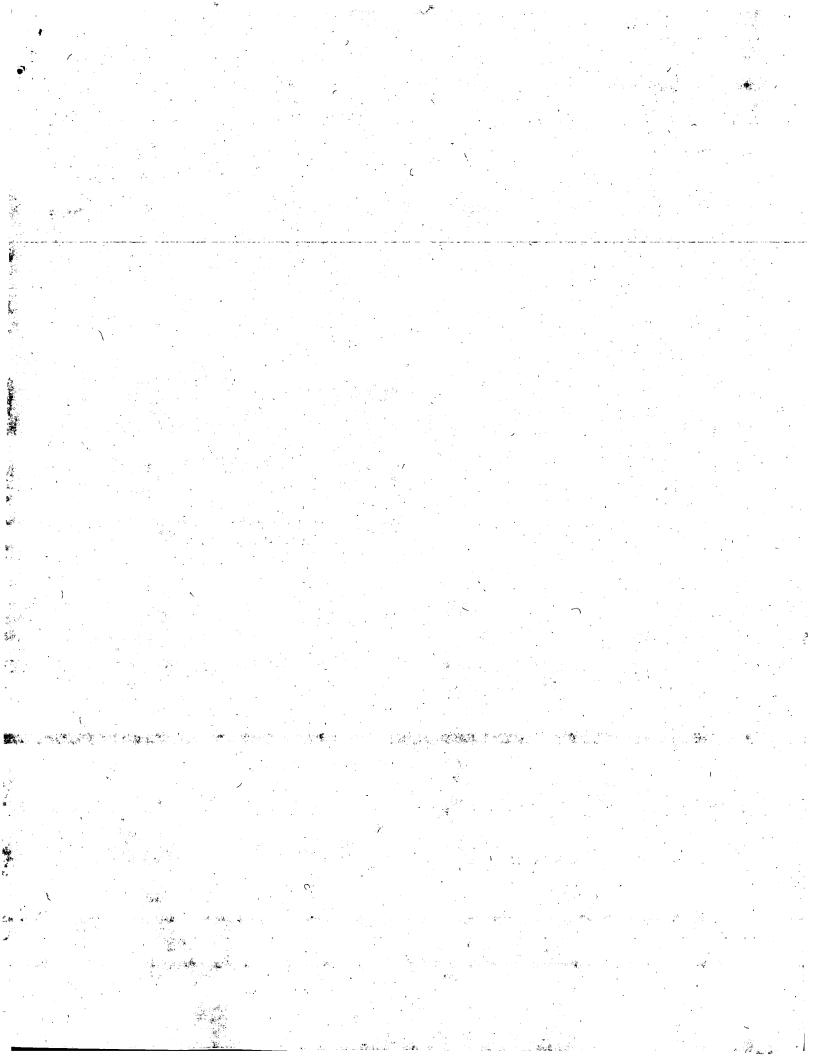
Ouery Match 60:0%; Score 36; DB 1; Length 1264;
Best Local Similarity 63:6%; Pred. No. 75;
Matches 7; Conservative 2; Mismatches 2; Indels 0; Gaps
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lery Match 60:0%; Score 36; DB 1; Length 1264;
sst Local Similarity 63.6%; Pred. No. 75;
itches 7; Conservative 2; Mismatches 2; Indels 0; Gaps
2 LQVLGLQLPTP 12
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Search completed: June 4, 2001, 12:24:09 Job time: 571 sec



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Minimum DB seq
Maximum DB seq
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Copyright (c) 1993 - 2000 Compugen Ltd.
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               Q9rz01 deinococcus
Q9tyt0 caenorhabdi
                                                            Q9nxb8 homo sapien
Q44982 caenorhabdi
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Q94010 caenorhabdi
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0910W7 0911t6 091860 0091860 004763 004763 095044 09504 09504 09504 09504 09504 09504 09504 099706 099706 09109 091099 099795 091099 099795 099795 099795 099795 099795 099795 099795 099795 099795	14	2	ۍ.	ъ	Ħ	G	4	4	ഗ	J.	10	10	10	4	N	w	2	4	10	4	10	10	14	11	Ħ	u
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# ALIGNMENTS

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RESULT 2
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Best Local Similarity
Matches 12; Conser
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Q92663;
Q1-FEB-1997
                                                                                                                                                                                                                              "Novel Fc gamma receptor I family gene products in human mononuclear cells.";
J. Clin. Invest. 90:2102-2109(1992).
EMBL; L03418; AAA36049.1;
HSSP; P12319; 1ALT.
INVERPRO; IPRO03006;
PFAM; PE00047; 1g; 3.
PRODOM; PD002534; -: 1.
SEQUENCE 374 AA; 42632 MW; D33D59398CEEA699 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-FEB-1997 (TrEMBLrel. 02, 01-FEB-1997 (TrEMBLrel. 02, 01-OCT-2000 (TrEMBLrel. 15,
                                                                                                                                                                                                                                                                                                                                                                                    Kimberly R.P.;
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                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=93055454;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                         277 ELQVLGLQLPTP 288
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PRELIMINARY;
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cha P.B., Doebele R.,
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Pred. No. 0.0034;
Mismatches 0
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          PRT;
          375 AA
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RESULT
Q92638
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Best Local S
Matches 11
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Mammalia; Euther
NCBI_TaxID=9606;
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01-FEB-1997 (TrEMBLrel. 0:
01-FEB-1997 (TREMBLrel. 0:
01-CCT-2000 (TREMBLREL. 1
01-CCT-2000 TREMBLREL. 1
              Porges A.J., Redecha P.B.,
Kimberly R.P.;
                                                                                                                           01-FEB-1997
01-OCT-2000
                               MEDLINE=93055454; PubMed=1430234;
                                                 SEQUENCE FROM N.A.
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                                                                                               Homo sapiens (Human)
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EMBL;
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EMBL;
EMBL;
    Novel Fc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Definition of interferon gamma-response elements in a novel human Fc gamma_receptor gene (Fc gamma RIb) and characterization of the gene
                                                                                                                                                                                                                                                                                                                        INTERPRO; IPR003006;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE OF 1-224 FROM N.A. MEDLINE=93018827; PubMed=1402657;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=BLOOD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                l Similarity 91.
                                                                                                                   RECEPTOR
                                                                           Eutheria;
    gamma receptor I
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7 (TrEMBLrel.
) (TrEMBLrel.
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                                                                                                                                                             PRELIMINARY;
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AAD13842.1;
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42881 MW;
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Primates;
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91.7%;
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family gene products in human mononuclear
                     Doebele R.,
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                                                                        Craniata; Vertebrata; Catarrhini; Hominidae;
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Best Local
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Homo. sapiens (Human).
Howo.sapiens (Human).
Therefore, Chordata; C.
Therefore, Primates; C.
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                                   Mammalia; Eutheria;
Bovidae; Bovinae; Bo
                                                    Bos taurus (Bovine)
Eukaryota; Metazoa;
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FC GAMMA RECEPTOR I.
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HSSP; P12319; 1ALT
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EMBL; L03420; AAA35826
HSSP; P12319; 1ALT.
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AAA35825.1; -.
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                                            Cetartiodactyla;
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Pred. No. 8.8;
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actyla; Ruminantia; Pe
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SEQUENCE
                                       Anopheles gambiae (African malaria mosquito).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Nematocera; CulicCulicidae; Anopheles.
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MEDLINE-99296679; PubMed=10366528;
ROOKER M.M., Sherburne C., Lawley T.D., Taylor D.
"Characterization of the Tra2 region of the IncHI
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Culicidae; Anopheles.
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Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Inse
Pterygota; Neoptera; Endopterygota; Diptera; Nematocera;
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Insect Biochem. Mol. Biol. 30:35-46(2000).
EMBL; AF117751; AAD38337.3; -.
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                               Eukaryota; Metazoa; Nemato Rhabditidae; Peloderinae;
                                                    Caenorhabditis elegans
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hemolymph in the mosquito Anop
Proc. Natl. Acad. Sci. U.S.A.
EMBL; AJ276428; CAB81934.1;
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Best Local
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                                                                                                                                                                                                                                                                                                               MOOD D.H., Nagdi M.A., Nascimento A.LT. O., Netto L.E.S.,
Nhani A. Jr., Nobrega F.G., Nunesi L.R., Oliveira M.A.,
de Oliveira M.C., de Oliveira R.C., Palmieri D.A., Paris A.,
Peixoto B.R., Pereira G.A.G., Pereira H.A. Jr., Pesquero J.B.,
Quaggio R.B., Roberto P.G., Rodrigues V., de Rosa A.J.M.,
de Rosa V.E. Jr., de Sa R.G., Santelli R.V., Sawasaki H.E.,
da Silva A.C.R., da Silva A.M., da Silva F.R., Silva W.A. Jr.,
da Silveira J.F., Silvestri M.L.Z., Siqueira W.J., de Souza A.A.,
vallada H., Van Sluys M.A., Verjovski-Almeida S., Vettore A.L.,
Zago M.A., Zatz M., Meddanis J., Setubal J.C.;
"The Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Common Com
                                                                                                                                                                                         Zago M.A., Zatz M., Meruanto V., The genome sequence of the plant pathogen Nature 406:151-157(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Coutinho L.L., Cristofani M., Dias-Neto E., Docena C., El-Dorry H., Facincani A.P., Ferreira A.J.S., Feireira V.C.A., Ferro J.A., Fraga J.S., Franca S.C., Franco M.C., Frobme M., Furlan L.R., Garnier M., Goldman G.H., Goldman M.H.S., Gomes S.L., Gruber A., Ho P.L., Hoheisel J.D., Junqueira M.L., Kemper E.L., Kitajima J.P., Krieger J.E., Kuramae E.E., Laigret F., Lambais M.R., Leite L.C.C., Machado M.A., Madeira A.M.B.N., Madeira H.M.F., Marino C.L., Marques M.V., Martins E.A.L., Martins E.M.F., Marino C.L., Marques M.V., Martins E.A.L., Martins E.M.F., Matsukuma A.Y., Menck C.F.M., Miracca E.C., Miyaki C.Y., Monteiro-Vitorello C.B., Monteiro, Nagai M.A., Nascimento A.L.T.O., Netto L.E.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-2035711, Simpson A.J.G., Reinach F.C., Arrida P., Abreu F.A., Acentro I.S., Simpson A.J.G., Reinach F.C., Araya J.E., Baia G.S., Baptista C.S., Alvarenga R., Alves L.M.C., Araya J.E., Baia G.S., Baptista C.S., Barros M.H., Bonaccorsi E.D., Bordin S., Bove J.M., Briones M.R.S., Barros M.H., Bonaccorsi E.D., Bordin S., Bove J.M., Briones M.R.S., Barros M.H., Camargo A.A., Camargo L.E.A., Carraro D.M., Carrer H., Colauto N.B., Colombo C., Costa F.F., Costa M.C.R., Costa-Neto C.M., Colauto N.B., Colombo C., Costa F.F., Costa M.C.R., Costa-Neto C.M., Colauto N.B., Colombo C., Costa F.F., Costa M.C.R., Costa-Neto C.M., Colauto N.B., Colombo C., Costa F.F., Costa M.C.R., Costa-Neto C.M., Colombo C., Costa F.F., Costa M.C.R., Costa-Neto C.M., Costa M.C.R., Costa
                                                                                                                                                            INTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=2371;
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01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29PB61;
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Science 282:2012-2018(1998).
EMBL; AL110477; CAB76742.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                222 KLOTLGIQLP 231
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[2]
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                                                                                                           PRO; IPRO01279; PF00753; lactar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Proteobacteria;
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                                                                                                 lactamase_B;
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70.0%;
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         MW;
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Pred. No.
A99CE5F8A6B557B9
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                                                                                                                                                                                                                                                                        Xylella fastidiosa."
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CRC64;
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Query Match

Score

38;

DВ

Length 317;

73060 MW;

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RESULT
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                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 7
                                       Kawakami T., Nuyuu.
Kawakami T., Mizuno T., Makajima Y., Mizuno T., Mizuno T., Mischi T., Shibahara T., J
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Matches 6; Conserv
                                                                                                                          Q9NXB8;
Q9NXB8;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-OCT-2000 (TrEMBLrel. 15, Last annotation updat
CDNA FLJ20335 FIS, CLONE HEP11429 (FRAGMENT).
 EMBL; AK000342; BAA91097.1;
NON_TER 649 649
SEQUENCE 649 AA; 73060 My
                              "NEDO human cDNA sequencing Submitted (FEB-2000) to the
                                                                                                     Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                        8EXN60
                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                          PROSITE; PS01212; P2X_RECEPTOR; PRODOM; PD002383; -; 1.
                                                                                                                                                                                                                                                                                                                                                                                                PRINTS; PR01307; P2XRECEPTOR. PRINTS; PR01309; P2X2RECEPTOR.
                                                                                                                                                                                                                                                                                                                                                                                                                   PFAM; PF00864; P2X_receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parker M.S., Larroque M
Hear. Res. 0:0,0(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                INTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                       INTERPRO; IPRO01429;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN-NIH STRAIN 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cavia porcellus (Guinea pig)
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01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=10141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P2X2 RECEPTOR
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                                                                                                                                                                                                                                                          QAKLLGLQNPTP
                                                                                                                                                                                                                                                                                                                                                                 channel;
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(TremBLrel. 07, Last sequence update)
)(TremBLrel. 15, Last annotation update)
"" "DITCE VARIANT P2X2-3.
                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Larroque M.L.,
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                                                                                                                                                                                                                                                                                                                                                     Transmembrane; AA; 55771 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: INTEGRAL MEMBRANE PROTEIN. BELONGS TO THE P2X RECEPTOR FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION:
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                                                                                                                                                                                                                                                          461
                                                                                                               Chordata;
Primates;
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                                                Morinaga
Tanaka T.
                                                                        Itoh
                           aka T., Nakamura Y
project.";
EMBL/GenBank/DDBJ
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Pred. No. 58;
3; Mismatches
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                                                                        T.
                                                                                                             Catarrhini; Hominidae;
                                                                                                                         Craniata; Vertebrata; Euteleostomi;
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4; Mismatches
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2B87E8B9F3CAC244
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                                                           Shigeta K., Senba T.,
M., Ota T., Suzuki Y.,
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58;
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                                                        Matsumura K.
Obayashi M.,
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Query Match Best Local

63.3%;

Score 38; Pred. No.

76;

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Length 649;

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RESULT
084727
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Best Local Similarity
Matches 7; Conserv
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01-JUN-1998
01-NOV-1998
                                                                                                                                                                                                                                                                       EMBL; AF043698; AAB97562.1;
NON_TER 1 1
SEQUENCE 134 AA; 13720 M
                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. STRAIN-BRISTOL N2; Waterston R.;
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Bonfield J., Burton J., Connell M., Copsey T., Cooper J., Coulson A.
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,
Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken I., Sonora A., Saunders D., Shownkeen F.
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J.,
Thierry-Mieg J., Thomas K., Vaudin M., Vaughan K., Waterston R.,
Watson A., Weinstock L., Wilkinson-Sproat J., Wohldman P.;
                                     01-NOV-1998
01-OCT-2000
                                                              084727;
01-NOV-1998
                                                                                       084727
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Tin-Wollam A., Graves T.,
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Rhabditidae; Pelode
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Chlamydia trachomatis.
                           PHOSPHOGLYCERATE
                                                                                                                                                                                                                                                                                                               Submitted
                                                                                                                                                                                                                                                                                                                                                                                 Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Nature 368:32-38(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=94150718; PubMed=7906398;
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                                 (TrEMBLrel. 08, Created)
(TrEMBLrel. 08, Last sequence update)
(TrEMBLrel. 15, Last annotation update)
                                                                                         PRELIMINARY;
                                                                                                                                                                                                           Conservative
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                        MUTASE
                                                                                                                                                                                                                                                                          13720 MW;
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70.0%;
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Last sequence update)
Last annotation update)
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Pred. No.
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Best Local
                                                                                                   Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2000 (TrEMBLrel. 14, Last annotation update)
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PROSITE; PS00175; PG_MUTASE; UNKNOWN_1.
SEQUENCE 226 AA; 25785 MW; 82AE9C780D400B8D CRC64;
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Science 282:754-759(1998).
EMBL; AE001343; AAC68317.1;
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MEDLINE=99000809; PubMed=9784136;
                                                                                                                                                                                                                                PFAM; PF01614;
SEQUENCE 253
                                                                                                                                                                                                                                                                                                                                      EMBL; AE001862; AAF12212.
                                                                                                                                                                                                                                                                                                                                                            radiodurans R1.";
Science 286:1571-1577(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID=1299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteria; Thermus/Deinococcus
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Fraser C.M.;
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                                                                                                                                                                                                                                                                                                                 TGR;
                                                                                                                                                                                                                                                                                                                                                                                                              Genome Sequence of the Radioresistant Bacterium Deinococcus
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205
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LAALGVSLPTP
                                             LQVLGLQLPTP
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                                                                                                   Similarity 7; Conser
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                                                                                                     Conservative
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AA;
215
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                                                                                                                                                                                                                                MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDonald I., Utterback T., Zalewski C.,
Daly M.J., Minton K.W., Fleischmann R.D.,
Salzberg S., Smith H.O., Venter J.C.,
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Pred. No.
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Pred. No. 39;
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Search completed: June

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GenCore version Copyright (c) 1993 - 2000

n 4.5 Compugen Ltd.

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Title:
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G49881	W38417	W43410	G42851	G09391	G50427	G05034	Y24479	Y24478	G42852	G09392	G50428	· G05035	G14199	G37548	G06464	G50429	G14200	G05036	G37549	G06465	Y16106	G37550	G06466	G33208	B42371	G40829	G40830	G40831	G50154	G50155	G50156	W86195	R12428
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# ALIGNMENTS

RESULT W60561 Identifying peptide(s) binding specifically to protein target - by expressing on phage surface and testing for binding to immobilised oligo:peptide derived from the target, useful for, e.g. identifying Extracellular domain; CD64; identification; antibody; immunohistochemical; immunofluorescent analysis; detection, W60561 standard; specific antibodies WPI; 1998-240964/21. 08-OCT-1996; 07-OCT-1997; Synthetic cell transformation; mutation; anti; oncogene. Oligopeptide from extracellular domain of CD64 18-AUG-1998 W60561; De Kruif CA, (UYUT-). RIJKSUNIV UTRECHT. 16-APR-1998. WO9815833-A1 (first entry) Logtenberg T; 96EP-0202791 97WO-NL00557. peptide; 12 AA.

W97833 W97834

Human

Fc receptor Fc receptor

cancer assoc macrophage-s high affinit

Synthetic oligopeptides domain of CD64. They were

They were synthesised

derived from the extrace on 25 polyethylene rods

from the extracellular

Example 1; Page 29; 40pp; English

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PN XX PD PR PR PR PR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cc adjacent, non-overlapping sets. Peptide specific phages were isolated by caffinity binding to the CD64-covered rods. The specification describes a affinity binding to the CD64-covered rods. The specification describes a cc method for the identification of a peptide able to bind specifically to a cc surface of a replicable display package, synthesising oligopeptides cc derived from the target protein on a solid phase, and testing for binding cc between the peptide and oligopeptides. The method is used to screen large cc between the peptide and oligopeptides. The method is used to screen large cc peptide libraries, especially to detect antibodies, or their fragments, cf forms of the same protein, including bispecific antibodies that bind to cell markers or that can differentiate between different cc forms of the same protein, including bispecific antibodies that bind to cepitopes on different molecules. The genes/oligonucleotides that bend cc selected peptides can be isolated and used for recombinant production of immunofluorescent analysis, and also to detect cell transformation caused cc mutitation in fantiformanae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                               13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
Aruffo A,
                                                                                                               01-DEC-1992;
25-FEB-1988;
                             (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                             25-FEB-1988;
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                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                    Cell surface antigen; cloning; immunoselection, therapy; diagnosis; vector; FCRI; Fc receptor.
                                                                                                                                                                                                                            US5506126-A.
                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                  Human FcRI (cDNA clone p98 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R91439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              by mutation in (anti)oncogenes.
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1 vwfhvlfylavg 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; Protein; 344
Seed B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                             92US-0983647.
88US-0160416.
89US-0379076.
90US-0553759.
                                                                                                                                                           88US-0160416
                                                                                                                                                                                                                                                                                       Location/Qualifiers 58
                                                                                                                                                                                                                                                      /note= "amino acid
translated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA.
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Pred. No. 1.1e-05;
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                                                                                                                                                                                                                                                      58 is Leu
product"
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                                                                                                                                                                                                                                                                                                                                                                  receptor
                                                                                                                                                                                                                                                                      in
                                                                                                                                                                                                                                                                      clone
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N-PSDB; T14718.

WPI;

1996-200279/20

Misc-difference

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/note= 82 /note= 64 /note= 60. /note= /note= 55

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Location/Qualifiers

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Query Match
Best Local Similarity
Matches 12; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                The amino acid sequence (R91439) of human FCRI was detd. from a cDNA clone, p98 (T14718), obtd. from a cDNA library using an immunoselection cloning method. FCRI is a high affinity receptor for the Fc portion of IgG, normally located on cell surfaces of macrophages. 2 Other cDNA clones, p135 (T14717) and p90 (T1419), of the p98 product is truncated compared with those of the p135 and p90 products (see also R91438 and W00859). A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                                       Sequence
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of diagnostic
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                                                                                                                                                                                                                                                                                                                                                                       344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NA encoding cell surface antigen - useful for isolation and therapeutic proteins
                                                                  100.0%;
                                                   Score 68; DB 17;
Pred. No. 0.00038;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          English
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RESULT
Y96183
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                                 Macrophage; FCRI; cell surface antigen; human; immunoselection; panning; immunodiagnosis; diagnosis; immunotherapy; gene therapy; immune disorder; infection; asthma; immune-complex disease; amyloidosis; multiple sclerosis; parasitic disease; Fc receptor.
Homo sapiens
                                                                                                                                                                  Human macrophage-specific FcRI.
                                                                                                                                                                                                                 19-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                             Y96183 standard; Protein; 344 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                    289 vwfhvlfylavg 300
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0,

Conservative,

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Mismatches

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Gaps

0;

Length 344; Indels

0

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01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                                                                   physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and treatment of immune-mediated infections, diseases, and disorders in animals, including humans. These disorders include asthma,
                                                                                                                                                                                                                   The present sequence is that of a human macrophage specific FcRI, as deduced from cDNA clone p135 (see A50631), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
     ability to
                                                                                                                                                                                                                                                                                                               Example 10;
                                                                                                                                                                                                                                                                                                                                                              Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stamenkovic I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-OCT-1998;
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                                                                                                                                                                                                      based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GEHO ) GEN
                                     Immune-complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                        2000-586382/55
DB; A50631.
                  normally located on the cell surfaces of macrophages.
                                                                                                                                                                                                                                                                                                               Column
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268
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900S-0553759
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89US-0379076
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338
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333
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332
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213
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139
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134
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129
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                                                                                                                                                                                                                                                                                                             53-55; 75pp;
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                                                                                                                                                                                                                                                                                                             English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGC"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGT
                                                                                                                                                                                                                                                                                                                                                                immune-complex disease, parasitic
cause it to occur on
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Sequence

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RESULT
R91438
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                       13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
sequence.
be helpful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein of FCRI and a receptor ligand will be helpful to increase the potencies of antibodies in therapy.
               for the Fc portion of IGG, normally located on cell surfaces of macrophages. Another isolated cDNA clone, p90 (T14719), coded for a variant (W00859) of the FcRI sequence, and a third clone, p98 (T14718), coded for an FcRI (R91439) having a different C-terminal sequence. A fusion protein of FcRI and a receptor ligand will
                                                                                                              The amino acid sequence (R91438) of human FCRI was detd. from cDNA clone, p135 (T14717), obtd. from a cDNA library using an
                                                                                                                                                                                                            Cloning of cDNA encoding cell surface antigen - useful for isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cell surface antigen; cloning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human FcRI (cDNA clone p135 product).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R91438;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R91438 standard; Protein; 374 AA
                                                                                                                                                                                                                                            WPI; 1996-200279/20
N-PSDB; T14717.
                                                                                                                                                                                                                                                                                         Aruffo'A,
                                                                                                                                                                                                                                                                                                                      (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                        01-DEC-1992;
25-FEB-1988;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US5506126-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-OCT-1996
                                                                                                                                                          Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunoselection cloning method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity hes 12; Conserv
                                                                                                                                                                                             diagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4
                                                                                                                                                                                                                                                                                         Seed B;
 in increasing the potency of antibodies in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            344 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                      92US-0983647.
88US-0160416.
89US-0379076.
99US-0553759.
93US-0139273.
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                                                                                                                                                                                             therapeutic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vector; FcRI; Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'amino acid 58 is Val in translated product"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     translated
                                                                                                           obtd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 68;
Pred. No.
                                                                                                                                                                                           proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                             FcRI is a high affinity receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25 is Thr
products"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunotherapy;
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Query Match Best Local :

Length 374;

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                                           Query Match
Best Local Similarity
Matches 12; Conserv
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                                                                                                                  The amino acid sequence (W00859) of human\FCRI was detd. from a CDNA clone, p90 (T14719), obtd. from a CDNA library using an immunoselection cloning method. FCRI is a high affinity recept for the Fc portion of IgG, normally located on cell surfaces of macrophages. Other CDNA clones (see also T14717-18) coded for variants (R91438-39) of the FCRI sequence. A fusion protein of FCRI and a receptor ligand will be helpful in increasing the potency of antibodies in therapy.
                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992;
25-FEB-1988;
13-JUL-1989;
13-JUL-1990;
18-OCT-1993;
                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                             WPI; 1996-200279/20.
N-PSDB; T14719.
                                                                                                                                                                                                                                                                                                             Aruffo
   289
                                                                                                                                                                                                                     Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                 (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                       25-FEB-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                             09-APR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5506126-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cell surface antigen; cloning; immunoselection; therapy; diagnosis; vector; FCRI; FC receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human FcRI (cDNA clone p90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W00859;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VWFHVLFYLAVG 12
||||||||||
|Vwfhvlfylavg 300
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                                                                                                                                                                                                                                                                                                            A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ch 100.0%; Score 68; DB 17; Similarity 100.0%; Pred. No. 0.00042; 12; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                             Seed
                                                                                                  374 AA;
                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
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89US-0379076.
90US-0553759.
93US-0139273.
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                                                                                                                                                                                                                                                                                                                                                                                                                    88US-0160416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein;
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                                                    100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           374
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                                            0,
                                                    Score 68; DB 17;
Pred. No. 0.00042;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA
                                           Mismatches
                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunotherapy;
                                                             Length 374;
                                          Indels
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                                                                                                                                                                         receptor
                                          0;
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                                          0
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RESULT W80448

This is the amino acid sequence of human Fc receptor I (FCRI), and deduced from CDNA clone pi35 (see V63456) isolated using a rapid immunoselection cloning method from a cDNA library expressed in cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extracorporeal interleukin-2

cos

Example 10;

Column 53-54; 75pp; English

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PR PR XX
                                                                                                                                                                                13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
                                                                       New cloning vector and polylinker - based on efficient cloning and expression of mammalian human lymphocyte antigenic sequences
                                                                                                              WPI; 1998-609251/51
N-PSDB; V63456.
                                                                                                                                          Aruffo A,
                                                                                                                                                                                                             01-DEC-1992;
25-FEB-1988;
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                                                                                                                                                            (GEHO ) GEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cloning.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fc receptor I; FcRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W80448 standard;
                                                                                                                                          Seed B;
                                                                                                                                                             HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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, 99
                                                                                                                                                                                                            92US-0983647
88US-0160416
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                                                                                                                                                                                         89US-0379076
90US-0553759
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171
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159
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85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein; 374 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cell surface antigen; lymphocyte;
                                                                                                                                                                                                                                                                                               "encoded by GGG"
                                                                                                                                                                                                                                                                                                                  "encoded
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                                                                                                                                                                                                                                                                                                                                                                                           "encoded by
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                                                                               existing sequences for cDNA(s), especially
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Best Local Similarity
Matches 12; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Human Fc receptor
Misc-difference
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                                                                                                                                Misc-difference
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cloning.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen by in eukaryotic cells and selection of cells expressing the antigen by
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 immune-mediated infections, diseases,
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77
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256
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176
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100.0%; Pr
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A novel method for cloning cDNAs
                                                                                                                                                                                                                                                                                                                                                                                                                      cell surface antigen; lymphocyte; human;
                                                                                                                                         "encoded by CAA'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (see X07373)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-JUL-1989;
13-JUL-1990;
21-MAY-1997;
07-JUN-1999 (first entry)
                                                                                           W97834 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression libraries is based on transient expression of an antigen in eukaryotic cells and selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the following the second second substrate.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New cloning vector and polylinker - based on existing sequences for efficient cloning and expression of mammalian cDNA(s), especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-609251/51.
N-PSDB; X07372.
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25-FEB-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treatment of immune-mediated infections, diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the isolation and cloning of any protein which can be expressed transported to the cell surface membrane of a eukaryotic cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  were also obtained.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p98 (see X07373)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5830731-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               including humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                               374 AA;
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88US-0160416
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Pred. No. 0.00042;
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Fc receptor I; FcRI; cell surface antigen; lymphocyte; human;

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Human Fc receptor I.

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 from a single patient undergoing extracorporeal interleukin-2 induction therapy. Clones p135 (see V63456) and p90 (see X07372) were also obtained. A novel method for cloning cDNAs from mammalian expression libraries is based on transient expression of an antigen by in enhanced in enhanced cells and selection of cells expression the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation and cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. It has been used to clone genes (see V63442-33) encoding cell surface antigen services of the cell surface membrane of the cell surface and transported to clone genes (see V63442-33) encoding cell surface
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13-JUL-1990;
21-MAY-1997;
                                                                                                                                                          This is the amino acid sequence of human Fc receptor I (FcRI), as deduced from cDNA clone p98/X2 (see X07373) isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS
                                                                                                                                                                                                                           Example 10; Column 53-54; 75pp; English.
                                                                                                                                                                                                                                                         New cloning vector and polylinker - based on existing sequences freefficient cloning and expression of mammalian cDNA(s), especially human lymphocyte antigenic sequences
                                                                                                                                                                                                                                                                                                                             WPI; 1998-609251/51
N-PSDB; X07373.
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                                                                                                                                             The cDNA library was constructed from polyA RNA of cells
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89US-0379076.
90US-0553759.
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                                                                                                                                                                                                                                                                                                                                                                                             panning; immunodiagnosis; diagnosis; immunotherapy; gene th
immune disorder; infection; asthma; immune-complex disease;
amyloidosis; multiple scierosis; parasitic disease; Fc rece
                                                                                                                                                                                                                                                                                                                                             Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                             Macrophage; FcRI; cell surface antigen; human; immunoselection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human macrophage-specific FcRI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Y96134;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Y96134 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genes can be expressed in a prokaryotic or eukaryotic host cells to produce the encoded protein. The invention also provides high efficiency expression vectors (see V63441 and V63444) which allow the generation of very large mammalian expression libraries. The purified genes and proteins are useful for immunodiagnostic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  animals,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunotherapeutic applications,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 289 vwfhvlfylavg 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 VWFHVLFYLAVG 12
                                                                      -difference
                                                                                                                                                                                                                                                                                                                                                                           sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               including
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immune-mediated infections,
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9 51
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0.00042;
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                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy;
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The present sequence is that of a human macrophage specific FCRI, as deduced from cDNA clone pl35 (see A50592), which was isolated from a cDNA library expressed in COS cells using a novel method of the invention designed to isolate CSA nucleic acids. The method is
                        surfaces other than macrophages, is useful protein of FCRI and a receptor ligand will the potencies of antibodies in therapy.
                                       immune complex disease, amyloidosis, parasitic diseases or multiple sclerosis. FCRI is a high affinity receptor for the Fc portion of IGG, normally located on the cell surfaces of macrophages. The ability to interfere with such bonding, or to cause it to occur on surfaces other than macrophages, is useful in therapy. A fusion protein of FCRI and a receptor ligand will be helpful to increase
                                                                                                                                animals, including humans.
                                                                                                                                              treatment of immune-mediated infections,
                                                                                                                                                        proteins they encode, are useful for immunodiagnostic and immunotherapeutic applications, including the diagnosis and
                                                                                                                                                                                      physical selection of cells expressing the antigen by adhesion to (panning on) an antibody-coated substrate such as a culture dish. CSA nucleic acids isolated by the method of the invention, and the
                                                                                                                                                                                                                                   based upon transient expression of a CSA in eukaryotic cells and
                                                                                                                                                                                                                                                                                                                         Example 10;
                                                                                                                                                                                                                                                                                                                                                               Isolated nucleic acid molecule encoding the CD19 cell surface antigen, useful for immunodiagnosis and immunotherapy of immune-mediated infections or disorders, e.g. asthma, immune-complex disease, parasiti
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23-MAR-1990;
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DB; A50592.
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88US-0160416.
89US-0379076.
90US-0498809.
90US-0553759.
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                                                                                                                              These disorders include asthma,
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                                                                                                                                              diseases, and disorders in
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Y96226
                               Query Match
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                                                                                                                                                                                                                                                                                                                                                        receptor consist only of the extracellular portion of the receptor and are not glycosylated i.e. they do not have transmembrane domains or signal peptides. The recombinant proteins may be used in immunoassays to determine the immune status of patients with chronic diseases of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; high affinity Fc receptor; FcgammaRI; immunoglobulin; infection; immune response; CD64; monocyte; macrophage; neut eosinophil; HIV; IGG; immunosuppressive; antirheumatic; cyto antiinflammatory; AIDS; SLE; multiple myeloma; rheumatoid ar
                                                                                                                                                                                                              compositions containing recombinant proteins may be used to treat or prevent autoimmune diseases, allergies or tumours, especially AIDS, rheumatoid arthritis or MM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     monocytes and macrophages, but expression can also be induced on neutrophils and eosinophils. Upon Fc receptor activation, immune effector pathways are activated, leading to immune response. The present sequence may be modified to produce recombinant versions. The recombinant Fc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FcgammaRI is also known as CD64. Fc receptors play an important role in defending the body against infections. First, pathogens are opsonised by serum immunoglobulins. The resulting complex then binds
                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                  immune system, e.g. AIDS, systemic lupus erythematosus (SLE), multiple myeloma (MM), or rheumatoid arthritis. In addition, pharmaceutical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cells expressing Fc receptors. FcgammaRI molecules are expressed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is the human high affinity Fc receptor, FcgammaRI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel Fc receptor lacking transmembrane domains, a signal peptide, glycosylation, useful for diagnosing and treating immune disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN
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                                                                                                                                                      374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 26-28;
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                            100.0%;
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Score 68; DB 21;
Pred. No. 0.00042;
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                                                           Length 374;
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cytostatic;
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Score 68; I

DB 21; ).00042; es 0;

Length 374; Indels

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1 VWFHVLFYLAVG

Sequence

B43683 ID B4 XX

B43683 standard; Protein;

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08-FEB-2001 B43683;

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314 vwfhvlfylavg 325

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                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                        antagonists may be also be B44240 represent sequences
                                                                                                                                                                                                                                                                                         viral infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antithyroid; antiallergic; antibacterial; antiviral; dermatological;
neuroprotective; cardiant; thrombolytic; coagulant; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and cells the genes are expressed in. Example of activities
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated nucleic acids comprising sequences encoding peptides useful for treating or diagnosing e.g. cancer -
                                                                                                                                                                                                                        invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vasotropic; antipsoriatic and antiangiogenic. The polynucleotides and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antiasthmatic; antirheumatic; antiarthritic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immune disorder; haematopoietic cell disorder; auto-
allergic reaction; graft versus host disease; organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis; cytostatic;
antidiabetic; antiasth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; cancer associated gene; cancer antigen; detection; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vasotropic;
                                                            Local Similarity 100 hes 12; Conservative
VWFHVLFYLAVG 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to C78448 encode the human cancer associated proteins given in to B44239. The proteins can have activities based on the tissues
                                                                                                                                                                         399 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ruben SM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  proliferative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antipsoriatic; antiangiogenic; gene therapy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   thrombolytic; cardiovascular disorder; infection; disease; drug screening.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              associated protein sequence SEQ ID NO:1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99US-0124270.
                                                                                                                                                                                                                                                           s. The peptides, nucleotides, antibodies, agonists and be also be used in drug screens. C78449 to C78457 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neuroprotective; thrombolytic; coagulant; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              iasthmatic; antirheumatic; antiarthritic; antiviral;
antithyroid; antiallergic; antibacterial; cardiant;
                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               proliferative;
                                                                                                                                                                                                                                          used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vulnerary;
                                                              0;
                                                          Score 68; DB 21;
Pred. No. 0.00045;
); Mismatches 0;
                                                                                                                                                                                                                                   in drug screens. C78449 to C78457 and in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immunomodulator; antidiabetic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               autoimmune disorder;
                                                                                                      Length 399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               immunomodulator
                                                            Indels
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                                                          Gaps
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XAXAX
                                                                                                   RESULT 13
W86195
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R12428
                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 10; Conserv
                                  10-MAY-1999
                                                            W86195
                                                                                   W86195 standard; Protein; 377
                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                           regulating antibody production in vivo and in plasmaphoresis to remove immune complexes or pathological antibodies. Rabbit IgG bound to cells transfected with the chimeric Fc(gamma) receptor
                                                                                                                                                                                                                                                                                                                                              This chimeric sequence is one example of a hybrid Fc receptor of the invention. The hybrid Fc receptor retains the Ig binding functions of the constituent receptor types. It can be used for
                                                                                                                                                                                                                                                                                               See also Q12078.
                                                                                                                                                                                                                                                                                                                                                                                                      Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                              antibody prodn.
                                                                                                                                                                                                                                                                                                                                                                                                                                       New immunoglobulin binding hybrid Fc receptor molecules - used to control auto-immune diseases and allergic reactions and to regulate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; Q12079.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hogarth PM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYME-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-OCT-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9106570-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunoglobulin-binding molecule; Fc(gamma)RI; Fc(gamma)RII.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R12428;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R12428 standard; Protein; 410 AA.
                                                                                                                                                    304 vwfhilfylsvg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hybrid Fc(gamma)RII/I receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-AUG-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Region
                                                                                                                                                                             1 VWFHVLFYLAVG 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1991-164135/22
                                                                                                                                                                                                                                                                                                                                                                                                   fig 3B; 80pp; English
                                                                                                                                                                                                                                                                        410
                                                                                                                                                                                                       Conservative
                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MELBOURNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hulett MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        89AU-0007045.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90WO-AU00513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= Fc(gamma)RI D3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                     315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'label= Fc(gamma)RII domains
                                                                                                                                                                                                                   94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytoplasmic domains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ierino FL,
                                                                                                                                                                                                                  Score 64;
Pred. No.
                                                                                     AA
                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                          chimeric Fc(gamma)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       McKenzie IFC,
                                                                                                                                                                                                                  0.0022;
                                                                                                                                                                                                                              DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           transmembrane and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   d1 and
                                                                                                                                                                                                                              Length 410;
                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Osman
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                                                                                                                                                                                                    Gaps .
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Human Fc receptor

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RESULT 14
G50156
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Best Local Similarity
Watches 9; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Allen J, Am
Oquendo C,
                                                                                                                                                                                                                                                                                                                                                                                                                                       This polypeptide human Fc receptor I (FCRI). FCRI cDNAs (see W81213) were isolated using a rapid immunoselection cloning method from a cDNA library expressed in COS cells. The cDNA library was constructed from polyA RNA of cells from a single patient undergoing extrcorporeal interleukin-2 induction therapy. DNA sequence analysis revealed that the cDNAs encoded type I integral membrane proteins with 3 extracellular immunoglobulin domains. The invention provides a novel method for cloning cDNAs from mammallian expression libraries. This is based on translent expression of an antigen in euwkaryotic cells and physical selection of cells expressing the antigen by adhesion to an antibody-coated substrate. The method is useful for the isolation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-1988;
13-JUL-1989;
23-MAR-1990;
13-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cDNA encoding human CD40 antigen - useful for cloning cDNA encoding cell surface antigens, constructing cDNA libraries, expression vectors for expression in eukaryotic cells or their fragments
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-DEC-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Key
Protein
18-OCT-2000
                                       G50156;
                                                                           G50156 standard;
                                                                                                                                                                                                                                                                                                                                                                               and molecular cloning of any protein which can be expressed and transported to the cell surface membrane of a eukaryotic cell. CD40 cDNA (see V81198) is specifically claimed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-069813/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Column 55-56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5849898-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fc receptor I;
                                                                                                                                                                     1 VWFHVLFYLA 10
|||:||||||
292 vwfnvlfyla 301
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Seed B,
                                                                                                                                                                                                                                                                                                                                                 377 AA;
                                                                                                                                                                                                                                                 Conservative
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92US-0983647.
88US-0160416.
89US-0379076.
90US-0498809.
90US-0553759.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD36; cell surface antigen; human; cDNA library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0485447
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1..377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
                                                                         Protein;
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Simmons I
                                                                                                                                                                                                                                                                   75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "this sequence contains a considerable number of differences from the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    deduced from the DNA sequence given in the specification"
                                                                         243 AA.
                                                                                                                                                                                                                                                 Score 51; DB 20; Pred. No. 0.36; 1; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lauffer L;
cc I, Stengelin
                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                     Length 377;
                                                                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                 0
                                                   18-JUN-1999;
18-JUN-1999;
18-JUN-1999;
18-JUN-1999;
18-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
                                                                                                                                                                     18-JUN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-FEB-1999;
05-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana protein fragment SEQ ID NO: 63527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-1999
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                                                                                                                                                                                                         8-JUN-1999
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                                                                                         99US-0139457.
99US-0139458.
99US-0139459.
99US-0139460.
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99US-01:
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99US-0139455
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RESULT
G50155
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Best Local S
Matches 6
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28-OCT-1999;
28-OCT-1999;
29-OCT-1999;
    Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter; termination sequence.
                                       Arabidopsis
                                                                           G50155;
                                                                                       G50155 standard; Protein; 257
                                                         18-OCT-2000
                                                                                                                                    :|||:||| :
219 iwfhilfyffi 229
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26-OCT-1999
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                                                                                                            15
                                                                                                                                                   1 VWFHVLFYLAV 11
                                                                                                                                                                                                                                                                                                            -OCT-1999
                                                                                                                                                                    Similarity 54.96; Conservative
                                      thaliana
                                                       (first entry)
                                                                                                                                                                                                     99US-0160741
99US-0160767
99US-0160770
99US-0160815
99US-0160815
99US-0160815
99US-0160880
99US-0161404
99US-0161406
99US-0161406
99US-0161359
99US-0161359
99US-0161351
99US-0161920
99US-0161920
99US-0161920
99US-0161920
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99US-0159638.
99US-0159584.
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990S-0153758
990S-0154039
990S-015486
990S-0155139
990S-0155139
990S-0156596
990S-0156596
990S-0156586
990S-0157753
990S-0157753
990S-0158029
990S-0158029
990S-0158029
990S-0158029
990S-0159294
990S-0159294
990S-0159295
990S-0159295
990S-0159295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0152363
                                      protein fragment SEQ ID NO:
                                                                                                                                                                             72.1%;
                                                                                                                                                                     ω,
                                                                                                                                                                    Score 49; DB 2
Pred. No. 0.49;
3; Mismatches
                                                                                          AA
                                                                                                                                                                                     DB 21;
                                                                                                                                                                    ۲,
                                     63526
                                                                                                                                                                                    Length 243;
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Gaps

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11-AUG

-sne6 -sne6 -sne6

3-0148341. 3-0148565.

02-AUG-1 03-AUG-1 04-AUG-1 04-AUG-1 05-AUG-1

1999

990S-0145087 990S-0145089 990S-0145145 990S-0145218 990S-0145218 990S-0145218 990S-0145218 990S-0145918 990S-0145919 990S-0145386 990S-0146386 990S-0146386 990S-0147038 990S-0147038 990S-0147204 990S-0148319

27-JUL-1999; 27-JUL-1999; 27-JUL-1999; 28-JUL-1999; 02-AUG-1999; 02-AUG-1999;

21-JUL-1999; 21-JUL-1999; 21-JUL-1999; 20-JUL-1999; 20-JUL-1999; 20-JUL-1999;

99US-0145085 990S-0145086 990S-0145088 99US-0144814 19-JUL-1999; 19-JUL-1999; 19-JUL-1999;

99US-0144085. 99US-0144086. 99US-0144325. 99US-0144331. 99US-014433. 99US-0144333. 99US-0144333.

3-JUL-1999

99US-0142920. 99US-0142977. 99US-0143542.

-0143624

99US-0141842. 99US-0142154. 99US-0142055. 99US-0142390. 99US-0142803.

990S-0139817. 990S-0139899. 990S-0140353. 990S-0140354. 990S-0140695. 990S-0140823. 990S-0140991.

20-AUG-1999; 20-AUG-1999; 20-AUG-1999; 23-AUG-1999; 23-AUG-1999; 25-AUG-1999; 26-AUG-1999; 27-AUG-1999; 27-AUG-1999;

-1999; -1999;

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47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	47.1	48.5	48.5	48.5	48.5	48.5	48.5		48.5	48.5
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US-09-188-930-144	US-08-602-359A-33	US-09-234-613-33	US-08-933-750C-33	US-09-370-253-18	5200327-17	US-08-318-193-6	US-08-318-193-39	US-08-556-597-168	US-08-620-717A-9	US-09-184-223-6	US-08-590-554A-6	US-08-485-568A-6	US-08-989-299-12	US-08-460-114A-4	US-08-262-338A-4	US-08-532-795-27	US-08-532-795-21
Sequence 144, App	Sequence 33, Appl	Sequence 33, Appl	Sequence 33, Appl	Sequence 18, Appl	Patent No. 5200327	Sequence 6, Appli	Sequence 39, Appl	Sequence 168, App	Sequence 9, Appli	Sequence 6, Appli	Sequence 6, Appli	Sequence 6, Appli	Sequence 12, Appl	Sequence 4, Appli	Sequence 4, Appli	Sequence 27, Appl	Sequence 21, Appl

ALIGNMENTS

### TELEPHONE: 201-831-3244 TELEFAX: 201-831-3305 INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS: LENGTH: 453 amino acids TYPE: amino acid Sequence 5, App... No. 5512472 PRIOR APPLICATION DATA: APPLICATION UNMBER: US 08/107,347 FILING DATE: 16-AUG-1993 ATTORNEY/AGENT INFORMATION: NAME: GOIDON, Alan M. REGISTRATION: NUMBER: 30,637 REFERENCE/DOCKET NUMBER: 854-012 (3: TELECOMMUNICATION INFORMATION: TELEPHONE: 201-831-3244 MOLECULE TYPE: ORIGINAL SOURCE: COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy APPLICANT: Kirsch, Donald R. TITLE OF INVENTION: A DNA Sequence APPLICANT: Bard, Martin APPLICANT: Kirsch, Dona PUBLICATION INFORMATION CORRESPONDENCE ADDRESS: APPLICANT: TITLE OF INVENTION: CLASSIFICATION: 435 APPLICATION NUMBER: FILING DATE: 11-MAN ORGANISM: STRANDEDNESS: single TOPOLOGY: linear COUNTRY: UZIP: 07470 ADDRESSEE: INFORMATION: Wayne Application US/08439131A New Jersey One Cyanamid U.S.A. Shimanuki, M. Goebl, M. Lai, Margaret H. K. Schizosaccharomyces pombe American Cyanamid Company peptide Floppy disk Reductase US/08/439,131A 5 854-012 (32,141) Version #1.25 Encoding Sterol Delta

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JS-08-440-674-4
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Patent No. 5525496
                                                                                                                                                                                                              NFORMATION FOR SEQ ID NO:
                                                                      PUBLICATION INFORMATION
                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/107,347
FILING DATE: August 16, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Alan M. Gordon
                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" 1.
                                                                                                                              OLECULE TYPE:
                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-831-3244
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Word Processor CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: American Cyanamid Company
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            || :||:::||
217 WF-ILFFISVG 226
                                                                                                                                                                                                           CITY: Wayne STATE: New -
                                                         AUTHORS:
                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 306 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ITLE OF INVENTION:
                                                                                                              DESCRIPTION:
                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: FILING DATE: May 1
                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE: Fission Yeast sts1+ Gene Encodes a Protein TITLE: Similar to the Chicken Lamin B Receptor JOURNAL: Molecular Biology of the Cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PAGES: 263-273
DATE: 1992
                                                                                                                                                                                 ENGTH:
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                                                                                                                                                                   amino acid
Fission Yeast stsl+ Gene
Encodes a
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                                                      M. Shimanuki,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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VENTION: A DNA Sequence Encoding Sterol
                                                                               Schizosaccharomyces
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                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                    IBM PC
                                                                                                                                                                               residues
                                                                                                           polypeptide
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                                                                                                                                                     single
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                                                     M. Goebl,
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Pred. No. 7
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                                                   M. ,Yanagida,
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Best Local Similarity
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DATE: 1992
                                                                         TELEFAX: (415) 324-167
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/071,880
FILING DATE: June 1, 193
APPLICATION NUMBER: 07/891,301
FILING DATE: May 29, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Hand Dolezalova
RECISCREATION NUMBER: 20,510
                                                                                                                          REGISTRATION NUMBER: 30,518
REFERENCE/DOCKET NUMBER: 48
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-1677
                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: DOS SOFTWARE: Wordperfect CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
COMPUTER: PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 5
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                               STRANDEDNESS:
                    TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 WEHVLEYLAVG
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: United States of America
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PC
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                  linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PHILLIPS, MOORE, LEMPIO & FINLEY
5 Sherman Avenue, Suite 6
peptide
                                                                                                                                                                                                                                                                                                                       03-APR-1995
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                                                                                                              324-1678
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DIAGNOSIS AND KIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       POLYPEPTIDES BINDING ANTI-
CRYPTOSPORIDIUM ANTIBODIES, DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  54.5%;
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                                                                                                                                                                                                                                                                                                                                     US/08/415,751
                                                                                                                                                           480.19-2 (HHD)
                                                                                                                                                                                                                                                                                                                                                                                                                3.5 inch, 1.44 Kb storage
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Pred. No. 77;
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; ORGANISM:
; FEATURE:
; NAME/KEY:
; NAME/KEY:
US-08-415-751-36
RESULT 5
US-08-624-125-8
; Sequence 8, Application US/08624125
; Patent No. 5744341
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Best Local S
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                               NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 703-413-3000
                                                                                                                                                                                                                                                                                                                                                                                                                              TTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              267 IWFXTIXYMWIG 278
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                                                                                              70 WWPVLWWIALG 80
                                                                                                                                                                                                                                                         TOPOLOGY: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: UZIP: 22202
                                                                                                                                                                                                                                                                                                                                                                                                NAME: KELBER, STEVEN B. REGISTRATION NUMBER: 30,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 29-MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 1755 S.
CITY: ARLINGTON
                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                     2 WFHVLFYLAVG 12
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                                                                                                                                                                                                                                                                                                  LENGTH:
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                                                                                                                                                    Similarity
5; Conserva
                                                                                                                                                                                                                                                                                   amino acid
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                                                                                                                                                                                                                                                                                               162 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Floppy disk
                                                                                                                                                                                                                                          protein
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                                                                                                                                                                                                                                                                     single
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                                                                                                                                                                  50.0%;
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                                                                                                                                                   Score 34; DB Pred. No. 47; 5; Mismatches
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Pred. No. 74;
4; Mismatches
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                                                                                                                                                                                DB 1; Length 162;
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US-08-663-310-
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Best Local Similarity
Matches 5; Conser
                                                                                                                                                                                                                                                                     Sequence 4, Application US/08663310 Patent No. 5811273
                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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APPLICANT: CUNNIN
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NFORMATION FOR SEQ ID NO:
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS
                                                                                                                   NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                       APPLICANT:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                    ITLE OF INVENTION:
                                                                                                                                                                  ITLE OF INVENTION:
                                                                                                                                                                                ITLE OF INVENTION:
                                                                                                                                                                                              PPLICANT:
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: UZIP: 22202
                                            COUNTRY:
                                                             STATE:
                                                                                     STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: KELBER, STEVEN B. REGISTRATION NUMBER: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 29-MA
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COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Reli CURRENT APPLICATION DATA:

PatentIn Release #1√0, Version #1.30

APPLICATION NUMBER:

23-SEP-1996

US/08/663,:310

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                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO P
FILING DATE: 26-DEC-1994
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MEDIUM TYPE: Floppy disk
RIOR APPLICATION DATA:
                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.30
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
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                                                                                                                                                                       APPLICATION NUMBER: US/0
FILING DATE: 23-SEP-1996
                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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26-DEC-1994
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-235917
FILING DATE: 05-SEP-1994
                                                                                                                              PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 0
FILING DATE: 23-SEP-1996
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LENGTH: 162 amino acids
                             PRIOR APPLICATION DATA:
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APPLICATION NUMBER: JP 5 FILING DATE: 27-DEC-1993
                                                                                               APPLICATION NUMBER: WO PCT/JP94/02220 FILING DATE: 26-DEC-1994
                                                                                                                                                                                                   FILING DATE
                                                                                                                                                                                                                                                                                                                                    ZIP: 20007-5109
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                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: BENT, Stephen A REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                    E: Foley & Lardner
3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                     USA
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YOKOYAMA, Akihiro
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27-DEC-1993
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45.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA STRANDS USEFUL FOR THE SYNTHESIS OF
               JP 5-348737
                                                                                                                                               us 08/663,310
                                                                                                                                                                                                                 US/09/006,49:
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Pred. No. 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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; MOLECULE TYPE: protein US-09-006-491-4
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                          COUNTY:
ZIP: 20007-510,
ZIP: 20007-510,
COMPUTER READABLE FORM:
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
COMPUTER: PC-DOS/MS-DOS
COMPUTER: PC-DOS/MS-DOS
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SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
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FORMATION FOR SEQ ID NO:
                                              REFERENCE/DOCKET NUMBER: 49441/109
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                             APPLICATION NUMBER: JP 5-348737
FILING DATE: 27-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REGERENCE COCCUER
                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US OF
APPLICATION STATE: 23-SEP-1996
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                                                                                                                                                                                      FILING DATE: 05-SEP-1
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                (202)672-5399
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RESULT 10
US-09-335-919-4
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Best Local Similarity
"hes 5; Conserve
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MOLECULE TYPE: US-09-335-919-4
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Patent No. 615013
                                                                         TELEX: 904136
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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LENGTH: 162 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                       FILING DATE: 26-DEC-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 23-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (202)672-5300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                       NAME: BENT, Stephen A. REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 3000 K St
CITY: Washington
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                              TOPOLOGY:
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                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: JP 6-235917
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                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/335,919
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                                                            LENGTH:
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                                           amino acid
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3000 K Street, N.W., Suite 500
                                                          162 amino acids
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YOKOYAMA, Akihir
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Akihiro
                                                                                                                                                                                                                                                                                                                                                        WO PCT/JP94/02220
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Pred. No. 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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69 IWAPVLWWIALG 80

1 VWFHVLFYLAVG 12

-09-335-919-11

Application US/09335919

ENERAL INFORMATION:

KONDO, Keiji

Akihiro

No. 6150130ihiko

ent No.

Query Match
Best Local Similarity
Matches 5; Conserv

Conservative

Score 34; Pred. No. Mismatches

Length 162;

Indels

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Gaps.

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                                                                                                               ; MOLECULE TYPE: US-09-335-919-11
                                Best Local Similarity 45.5%;
Matches 5; Conservative
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                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 49
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
                                                                                                                                                       TYPE: amino acid
                                                                                                                                           TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: JP 5-348737 FILING DATE: 27-DEC-1993
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2 WFHVLFYLAVG 12
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                                                                                                                                             linear
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                                                       Score 34;
Pred. No.
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                                            Mismatches
                                                            DB
47;
                                                                        Length 162;
                                            Indels
                                             0;
                                            Gaps -
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GENERAL INFORMATION:

CORRESPONDENCE NUMBER OF SEQUENCES:

ADDRESS

APPLICANT: Murphy, Randall B.
APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF

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US-08-332-562A-133
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                               RESULT 13
US-08-118-270-49
Patent No. 5508384
                                                                                                                                                    Best Local Similarity 45...
Matches 5; Conservative
                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                              TOPOLOGY: 1
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                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 54
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 27-MAY-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
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NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
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                                                                                           211 VWYHTAFSLVM 221
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                         1 VWFHYLFYLAV 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       133, Application US/08332562A
o. 5985599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20007-5109
                                                                                                                                                                                                                                                                                         : 261 amino acids amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INVENTION:
               Application US/08118270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3000 K Street; N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HOGARTH, Mark P.
                                                                                                                                                                                                                                                                                                                                                                  (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HULETT, Mark D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BONADONNA, Lisa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIBBS, Margaret L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                              inear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                              protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bernadette M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FC RECEPTOR FOR IMMUNOGLOBULIN
                                                                                                                                                                   50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            us 07/896,457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US/08/332,562A
                                                                                                                                                                                                                                                                                                                                                                                                                              29,768
                                                                                                                                                                                                                                                                                                                                                                                                                 54270/119/GRHA
                                                                                                                                                                      Score 34; DE Pred. No. 79;
                                                                                                                                                           Mismatches
                                                                                                                                                                                      Length 261
                                                                                                                                                           0
                                                                                                                                                         Gaps
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SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/335,919

OPERATING SYSTEM:

IBM PC compatible PC-DOS/MS-DOS

Version #1.30

PRIOR APPLICATION DATA:

CLASSIFICATION: FILING DATE: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

20007-5109

CORRESPONDENCE ADDRESS

ADDRESSEE:

E: Foley & Lardner 3000 K Street, N.W., Suite 500

STREET: 3000 K St CITY: Washington

COUNTRY:

USA

NUMBER OF SEQUENCES: ITLE OF INVENTION:

TLE OF INVENTION:

XANTHOPHYLLS

DNA STRANDS USEFUL FOR THE SYNTHESIS OF XANTHOPHYLLS AND THE PROCESS FOR PRODUCING THE

XANTHOPHYLLS AND THE PROCESS

419 Seventh

Suite 300

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Best Local Similarity 04...
Ches. 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                  sequence 49, Application PC/TUS9308528 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08, FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07, FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 202-737-3528
TELEX: 248633
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                         CURRINT APPLICATION DATA:
APPLICATION NUMBER: PC
FILING DATE: 09-SEP-19
PRIOR APPLICATION DATA:
                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                        APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: BROWDY
STREET: 419 Sevent
CITY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                  ORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 VWYYGLFY 68
                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                  COUNTRY: U
ZIP: 20004
                                                                                                                                                                                                                                                               STATE: D.C.
                                                                                                                                                                                                                                                                               CITY: Washington
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
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REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 VWFHVLFY 8
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                                                                                                                                                                                                                                                                                             E: BROWDY AND NEIMARK 419 Seventh Street, N.W.,
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                                                                                                                                                                                                                                                                                                                                                                           New York University
VENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                USA
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                                                                                                                                             Patentin Release #1.0, Version #1.25
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                                                                                                             PCT/US93/08528
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Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                  Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 340;
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RESULT 15
US-07-937-609-20
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MOLECULE TYPE: peptide
PCT-US93-08528-49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 62.5
5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No.
                                                                                                  NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 4039
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                  INFORMATION FOR SEQ ID NO:
                             SEQUENCE CHARACTERISTICS:
LENGTH: 372 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/861,769
FILING DATE: 01-APR-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: WANK, Stephen A.
TITLE OF INVENTION: CLONING AND FUNCTIONAL EXPRESSION OF
TITLE OF INVENTION: CHOLECYSTOKININ RECEPTOR-ENCODING DN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                            APPLICATION NUMBER: US 07/928,033 FILING DATE: 11-AUG-1992 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/831,248 FILING DATE: 07-FEB-1992
                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 19
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 VWYYGLFY 68
                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: UPPLING DATE: 19920902
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TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 22313-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Alexandria
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Pred. No. 1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Version #1.25
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Search completed: June 4, 2001, 12:14:33 Job time: 226 sec	Db 97 VWYYGLEY 104	Matches 5; Conservative Qy 1 VWFHVLFY-8	Query Match Best Local Similarity	US-07-937-609-20
4, 2001,		ative	50.0%; 62.5%;	•
12:14:		2; Mis	Score Pred.	
3 3		2; Mismatches 1; Indels	50.0%; Score 34; DB 1; Length 372; 62.5%; Pred. No. 1:2e+02;	

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Result
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Maximum Match 100%
Listing first 45 summaries
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BG
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length:
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1: pir1:
2: pir2:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
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pir3:*
pir4:*
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ATPase subunit 6 -
                             Fc-gamma receptor
Fc-gamma RIIIB-alp
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52.9	52.9	52.9	52.9	52.9	52.9	52.9	52.9	52.9	52.9	52.9	54.4	54.4	54.4	54.4	54.4
555	515	483	370	368	366	254	252	. 242	123	82	. 776	621	473	407	405
2	Ν	N	1	N	۲	N	N	N	2	N	N	N	N	N	N
нвзо43	H84013	T16443	XNVKUD	I52863	XNBYUG	T20882	G75485	E81315	S63055	T51909 · ·	148317	T48646	S51256	н81160	E81946
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### ALIGNMENTS

C;Species: Homo sapiens (man)
C;Date: 02-Jul-1996 #sequence\_revision

C; Accession: I55577;

N; Contains: Fc-gamma (IgG)

Fc gamma (IgG) receptor I-B splice form 1 precursor - N.Alternate names: CD64

receptor I-B splice form

02-Jul-1996 #text\_change 21-Jan-2000

human

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Fc gamma (IgG) receptor I (high affinity) form b - human N;Alternate names: CD64 C;Species: Homo sapiens (man) C;Date: 03-Apr-1992 #sequence_revision 03-Apr-1992 #text_change 21-Jan-2000 C;Recession: A41357; S03019
                                                                                                                                                           RESULT
A41357
                                                                                                                                                                                                                                                                   밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Map position: 1p12-1p12
C;Superfamily: Fc gamma receptor I; immunoglobulin homology
C;Superfamily: Fc gamma receptor; immunoglobulin receptor; transmembran
C;Reywords: alternative splicing; glycoprotein; immunoglobulin receptor; transmembran
F;117-170/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Gene: GDB:FCGR1B; CD64
A;Cross-references: GDB:135923; OMIM:601502
R;Allen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Experimental source: A; Note: splice form B2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Porges, A.J.; Redecha, P.B.; Doebele, R.; Pan, L.C.; Salmon, J.E.; Kimberly, R.P.
J. Clin: Invest. 90, 2102–2109, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Reference number: 155577;
A;Accession: 155577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: mRNA
A; Residues: 1-280 < RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Title: Novel Fc gamma
                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;Cross-references: GB:L03419; NID:g182460; PIDN:AAA35825.1; PID:g292023;Note: splice form B1;Accession: I70303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         \;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Comment: This receptor does not bind monomeric IgG with high affinity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cross-references: GB:L03420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Status:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Residues: 1-10,103-153,'A',155-280 <RE2>
                                                                                                                                                                                                                                                                195 VWFHVLFYLAVG 206
                                                                                                                                                                                                                                                                                                                                                                       y Match 100.0%; Score 68; D
Local Similarity 100.0%; Pred. No. 0.
hes 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                  1 VWFHVLFYLAVG 12
J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 :L03420; NID:g182461; mononuclear cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         eptor I family gene products in human mononuclear cells. MUID:93055454
                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
0.00015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PIDN:AAA35826.1; PID:g292024
                                                                                                                                                                                                                                                                                                                                                                                                                      Length 280;
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                                                                                                                                                                                                                                                                                                                                                                       Gaps
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R; Perez,
                                                                                                                                               A; Title: Nucleotide sequence of three cDNAs A; Reference number: S03018; MUID:89098339
                                                                                                                                                                                         R;Allen, J.M.; Seed, B.
Nucleic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N; Alternate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      F;117-170/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Title: Nucleotide sequence of three cDNAs
A;Reference number: S03018; MUID:89098339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic Acids Res. 16, 11824, 1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Allen, J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: GB:X14355; GB:M21090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;Porges,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; van de Winkel,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cross-references: EMBL:X14355; NID:g31333; PIDN:CAA32536.1; PID:g31334 ...;Note: the authors translated the codon ACT for residue 25 as Ala :Superfamily: Fc gamma receptor I; immunoglobulin homology :Superfamily: Expansarive splicing; immunoglobulin receptor; transmembrane protein; Keywords: alternative splicing; immunoglobulin receptor; transmembrane protein; 117-170/Domain: immunoglobulin homology <IMM>
                                              Molecule type: mRNA
Residues: 1-24, 'S',26-337, 'T',339-374 <AL2>
Cross references: BMBL:X14356; NID:931331; PIDN:CAA32537.1;
Perez, C.; Wietzerbin, J.; Benech, P.D.
                                                                                                                                                                                                                          Molecule type: mRNA
Residues: 1-24,'S',26-337,'T',339-374 <AL1>
Cross-references: GB:X14356; GB:M21091; NID:g31331;
                                                                                                                                                                                                                                                                                            Reference number: Accession: B41357
                                                                                                                                                                                                                                                                                                                                                         Allen, J.M
Pience 243,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       . Biol. Chem. 266, 13449-13455, 1 Title: Gene organization of the Reference number: A39878; MUID:9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reference number: I55577; MUID:93055454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Title: Novel Fc gamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross-references: GB:M63830; GB:M63835; NID:g180277; Porges, A.J.; Redecha, P.B.; Doebele, R.; Pan, L.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Date: 30-Dec-1991 #sequence_revision 06-Sep-1996 #text_change Accession: A39878; I70304; B41357; S03018; I57525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Residues: 1-344 <ALL2>
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                            Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                  Residues: 1-374 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Status: translated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Accession: A39878
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Accession:
                                                                                                                                ccession: S03018
                                                                                                                                                                                                                                                                                                                                                                                                                                    Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ccession:
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          Two
                                                                                                                                                                                                                                                                                                            Isolation and expression of functional nce number: A41357; MUID:89100284
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                                                                                                                                                                                                                                                                                                                                                                            J.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Invest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IgG) receptor I-A (high affinity) precursor
                               Biol. 13,
cis-DNA elements involved in myeloid-cell-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1-344 <ALL1>
                                                                                                                                                                                                                                                                                                                                                    378-381,
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12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            I70304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -374 <VAN>
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                                                                                                                                                                                                                                                                                                                                                                            Seed, B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90, 2102-2109, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J.G.J.; Ernst, L.K.; Anderson,
                                                                                                                                                                                                                                                                                                                                                                                        GB:L03418; NID:g184840;
                               2182-2192,
                                                                                                                                                                                                                                                                                                                                                       1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the human high affinity receptor MUID:91302383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              family
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NID: 931333; PIDN: CAA32536.1;
                                                                                                                                                                      for
                                                                                                                                                                                                                                                                                                                                                                                        PIDN: AAA36049.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for
                                                                                                                                                                      the
                                                                                                                                                                                                                                                                                                                             high-affinity
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                                                                                                                                                                    human
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                                                                                                                                                                                                                             PIDN:CAA32537.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chiu,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PIDN:AAA35678.1; PID:gli
Salmon, J.E.; Kimberly,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in human mononuclear cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human
                                                                                                                                                               high affinity Fc receptor
                                                                                                                                                                                                                                                                                                                                                                                        PID:g292169
                                                                 PID:g31332
                                                                                                                                                                                                                                                                                                                             Fc receptor complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for
    expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-Jul-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IgG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                        PID:g31332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PID: 931334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PID:g180279
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      and
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                                                                                                                                                               (FC
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                                            A;Status:
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                                                                                                                                                                                                                                                                  A46480
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                                                                                                                                                                                                                                                                                     RESULT
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A;Title: Linkage on chromosome 3 of autoimmune diabetes and A;Reference number: I48471; MUID:93242399
A;Accession: I48471
                                                                                                                                                                          Fc gamma (IgG) receptor high affinity - mouse
C:Species: Mus musculus (house mouse)
C:Date: 02-Jul-1996 #sequence_revision 02-Jul
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Theres 12; Conserv
                                                                                                                  R;Prins, J.B.; Todd, J.A.; Rodrigues, N.R.; Ghosh, S.; Science 260, 695-698, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        F;117-170/Domain: immunoglobulin homology <IMM2> F;293-313/Domain: transmembrane \#status predicte
              A; Molecule type: mRNA
                                      A; Status: preliminary; translated
                                                                                                                                                                                                                    Fc gamma (IgG)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       F;59,78;152,159,163,195,240/Binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C; Superfamily: Fc gamma receptor I; immunoglobulin homology C; Roywords: alternative splicing; glycoprotein; immunoglobu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Gene: GDB: FCGR1A; CD64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Cross-references:
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A; Residues: 1-8 < RE2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: 157525; MUID: 93204964
                                                                                                                                                           Accession: I48471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                        289 VWFHVLFYLAVG 300
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92/Domain: artracallular #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                               1 VWFHVLFYLAVG 12
1-336 <RES>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1-8 <RE2>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GB:S57204; NID:g298692;
                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score
100.0%; Pred.
                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                    from GB/EMBL/DDB
                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 site: carhohuz
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                          02-Jul-1996 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                              68;
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                                                                                                                                                                                                                                                                                                                                                                                                                        0.0002
                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 374;
                                                                                                                                   Hogarth,
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                                                                                             defective
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                                                                                                                                   P.M.; Wicker,
                                                                                                                                                                            21-Jan-2000
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                                                                                           Fc receptor
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                                                                                                                                 L.S.; G
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Fc gamma (IGG) receptor high affinity - mouse N;Alternate names: high affinity IGG receptor C;Species: Mus musculus (house mouse) C;Date: 18-Jun-1993 #sequence_revision 18-Nov-1994 #text_change 21-Jan-2000 C;Accession: A46480; A43511 R;Osman, N.; Kozak, C.A.; McKenzie, I.F.; Hogarth, P.M. J. Immunol. 148, 1570-1575, 1992
                                  A; Reference number: A46480; A; Accession: A46480
                                                                 A; Title: Structure and mapping
                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                   303
                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                1 VWFHVLFYLAVG 12
               preliminary
type: DNA
                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                              94.1%;
                                                 MUID:92166399
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Pred.
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No.
                                                               encoding
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0.0009;
                                                                 mouse
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                                                            high affinity
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                                                                                                                                                                                                                                                                                                                           Gaps
                                                               Fc
                                                               gamma
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RΙ

homology

PID: g311749

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hypothetical protein TM0089 - Thermotoga maritima (strain MSB8)
C;Species: Thermotoga maritima
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 21-Jul-2000
                                                                                                                        RESULT
C72420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-404 <SEA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Note: sequence extracted from R; Sears, D.W.; Osman, N.; Tate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Science 277, 1453-1462, 1997
A; Title: The complete genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probable membrane protein yick - Escherichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: GB:M31314; NID:g200752; PIDN:AAA40056.1; C;Superfamily: Fc gamma receptor I; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Fille: Molecular cloning and expression of the mouse high affinity Fc receptor A;Reference number: A43511; MUID:90111035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     J. Immunol. 144, 371-378, 1990 A; Title: Molecular cloning and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Accession: A43511
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;TILLE: The complete genome sequence of Escherichia;Reference number: A64720; MUID:97426617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: GB:AE000443; GB:U00096; NID:g2367255; Experimental source: strain K-12, substrain MG1655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;Blattner, F.R.; Plunkett III,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                         ;375-391/Domain:--transmembrane
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;Accession: D65167
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                                                                                                                                                                                                                                                                                                        Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Molecule type:
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Best Local Similarity
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105-121/Domain: transmembrane #status predicted <TM4>
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                                                                                                                                                                                                                                                                                                                                                                                                              4-330/Domain: transmembrane
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                                                                                                                                                                                                                                             1 VW-FHVLFYLAVG 12
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K.E.; Clayton, R.A.; Gill, S.R.; Gwinn, M.L.; Dodson, R.;
M.M.; Stewart, A.M.; Cotton, M.D.; Pratt, M.S.; Phillips,
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                                        C72420
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53.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI backbone (NCBIN:85205, NCBIN:85208, B.; McKenzie, I.F.C.; Hogarth, P.M.
                                                                                                                                                                                                                                                                                                                                                                                         #status
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#status predicted
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Pred. No. 3.8;
4; Mismatches
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Pred. No. 0.0011;
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R.J.; Haft,
ps, C.A.; Ri
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Hickey
on, D.,
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B
                               A;Molecule type: DNA
A;Residues: 1-1852 <FUJ>
A;Cross-references: DDBJ:AB000125; NID:g2308976; PID:d1022569; PID:g2308977
A;Cross-references: DBJ:Amembrane-bound protein. It is involved in the co
                                                                                                                                                                               R; Fujiwara, M.; Horiuchi, H.; Ohta, A.; Takagi, M. Biochem. Biophys. Res. Commun. 236, 75-78, 1997 A; Title: A novel fungal gene encoding chitin synth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Evidence for lateral gene transfer between Archaea and A;Reference number: A72200; MUID:99287316
A;Accession: C72420
                   C;Comment: This enzyme is a membrane-bound
C;Genetics:
                                                                                                                                                                                                                                                                                 chitin synthase (EC 2.4.1.16) asmA - Emericella nidulans N;Alternate names: chitin-UDP N-acetylglucosaminyltransferase C;Species: Emericella nidulans, Aspergillus nidulans
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C;Superfamily: Thermotoga maritima hypothetical protein TM0089
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                                                                                                                    A; Status: nucleic acid sequence not shown
                                                                                                                                       A; Reference number: JC5546; MUID: 97366599
A; Accession: JC5546
                                                                                                                                                                                                                                             C; Accession:
                                                                                                                                                                                                                                                                C;Date: 02-Sep-1997 #sequence_revision 05-Sep-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Map position:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Gene: CESP:C50Cl0.4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:Z72505; PIDN:CAA96610.1; GSPDB:GN00023; CESP:C50C10.4 A;Experimental source: clone C50C10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
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Best Local Similarity 54...
6; Conservative
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Best Local
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8; Conser
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54.5%;
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Pred. No. 8.
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Pred. No. 4.1;
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A;Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, S.; Galizzi, A lech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Hosono, S.; Hu Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.; Lapidus, A.; La, Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; Masuda, S., M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl, T.M.; P. Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Y.; Sato, T.; A; Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, J.; Sekowska, akeuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.; Tosato, V.; T.; Winters, P.; Wipat, A.; Yamamoto, H.; Yamane, K.; Yasumoto, K.; Yata, K.; Yasumotos: Yoshikawa, H.F.; Zumstein, E.; Yoshikawa, H.; Danchin, A. A; Title: The complete genome sequence of the Gram-positive bacterium Bacillus s. A; Reference number: A69580; MUID:98044033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Date: 05-Dec-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hypothetical protein a70L - Chlor
C:Species: Chlorella virus PBCV-1
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F;102-110/Domain:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Accession: T17560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Molecule type: DNA
; Residues: 1-88 <GRA>
Cross-references: GB:Z99105; GB:AL009126; NID:g2632457; PIDN:CAB11990.1; PID:e1182148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Accession: A69747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cross-references: EMBL:U42580; Experimental source: specific )
                                                Molecule type: DNA
                                                                 Status: preliminary; nucleic acid sequence not shown;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Species:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
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Date: 15-Oct-1999 #sequence_revision 15-Oct-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Reference number: Z18806
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity hes 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                    ; Ogasawara, N.; Moszer, I; Albertini, A.M.; Alloni, G.; Azevedo, V.; .; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.I., S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 66. 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T17560
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: ATP-binding #status
                              <KUN>
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70.0%;
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host Chlorella strain NC64A
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Pred. No.
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3.5;
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                                                              translation not shown
                                                                                                                                                                                                                                                                                                                                                          S.; Galizzi, A.; Galle
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, C.; Ferrari,
                                                                                                                                                                                                 V.; Uchiyama
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1-421 <BLAT>

Escherichia

coli

hypothetical

protein (glnA-fdhE intergenic

region)

PIDN:AAC76871.1; PID:g17903

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A;Cross-references: EMBL:L19201; NID:g304961; PIDN:AAB03008.1; PID:g304979
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1993
R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; Riley, M.;
A.; Rose, D.J.; Mau, B.; Shao, Y.
                                           A; Title: The complete genome sequence of Escherichia coli K-12
A; Reference number: A64720; MUID:97426617
                         A; Accession: E65192
                                                                                        Science 277, 1453-1462,
                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-421 < PLU>
                                                                                                                                                                                                                                                                                                                                                                              C;Date: 19-May-1994 #sequence_revision 01-Sep-1995 #text_change 24-Sep-1999
C;Accession: S40819; E65192
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                                                                                                                                                                                                                                                                                        A; Reference number:
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                                                                                                                                                                                                                                                                                                                                                            R;Plunkett III,
                                                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical 46.3K protein (glna-fdhe intergenic region) - Escherichia coli
N;Alternate names: hypothetical protein o421
Status: preliminary; nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                             Status: preliminary; nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                      Accession: S40819
                                                                                                                                                                                                                                                                                                                                                                                                                            Species:
                                                                                                                                                                                                                                                                                                                 Title: Analysis of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
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nes 6; Conserv
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                                                                                                                                                                                                                                                                                        S40802;
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                                                                                                                                                                                                                                                                                        Escherichia coli
)2; MUID:93347969
                                                                                                           Shao,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  58.8%;
                                                                                                                                                                                                                                                                                                                                   V.; Daniels, D.L.; Blattner, 3398, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 40;
Pred. No.
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17;
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                                                                                                                                                                                                                                                                                                            III.
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fro

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Matches
                                         Query Match
Best Local
171
               1 VWFHVLFYLAVG 12
IWFYSLIVIAVG 182
                                Similarity 6; Conser
                                Conservative
                                       57.4%;
50.0%;
                                       Score Pred.
                                       39;
No.
                                       DB
24;
                                               Length 421;
                                Indels
                               0;
                              Gaps
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hypothetical protein MTH768 - Methanobacterium thermoautotrophicum (strain Delta C;Species: Methanobacterium thermoautotrophicum C;Date: 05-Dec-1997 #sequence\_revision 05-Dec-1997 #text\_change 22-Oct-1999 #sequence\_revision 05-Dec-1997 #text\_change 22-Oct-1999

H)

R;Smith, C; Accession: F69202 ;Smith, D.R.; Doucette-Stamm, L.A.; Deloughery, C.; Lee, H.; Dubois, J.; Aldredge, T. Qiu, D.; Spadafora, R.; Vicaire, R.; Wang, Y.; Wierzbowski, J.; Gibson, R.; Jiwani, S.; Church, G.M.; Daniels, C.J.; Mao, J.; Rice, P.; Noelling, J.; Reeve, J.N. , S.; Church, G.I Bacteriol. 179, G.M.; Daniels, C.J 79, 7135-7155, 1997

A;Reference number: A; Title: Complete genome sequence A69000; quence of Methanobacterium thermoautotrophicum Delta MUID:98037514

A; Molecule A; Accession: F69202 A;Status: preliminary; type: DNA: 1-84 <MTH> nucleic acid sequence not shown; translation

A; Experimental source: C; Genetics: A; Residues: GB:AE000855; GB:AE000666; NID:g2621852; be: strain Delta H PIDN: AAB85271.1; PID:9262

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В
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Best Local Similarity
Thes 4; Conserve
                                                                                                  submitted to the EMBL Data Library, November 1993 A; Description: Mycobacterium leprae cosmid B1496. A; Reference number: S72693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MLCL536.34 protein -
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A,Authors: Foulger, D.; Fritz, C.; Fujita, M.; Fujita, Y.; Fuma, A,Authors: Foulger, D.; Henaut, A.; Hilbert, H.; Holsappel, S.; lech, J.; Harwood, C.R.; Henaut, A.; Hilbert, H.; Holsappel, S.; Koetter, P.; Koningstein, G.; Krogh, S.; Kumano, M.; Kurita, K.;
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                                                                                                                                                                                                                                                                                                                                                                   ;Parkhill, J.; Barrell, B.G.; Rajandream, M.A. ubmitted to the EMBL Data Library, September 1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rieger, M.; Rivolta, C.; Rocha, E.; Roche, B.; Rose, M.; Sadaie, Authors: Schleich, S.; Schroeter, R.; Scoffone, F.; Sekiguchi, keuchi, M.; Tamakoshi, A.; Tanaka, T.; Terpstra, P.; Tognoni, A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Authors: Lauber,
                                                                                  Accession:
                                                                                                                                                                                 Cross-references: EMBL:299125; NID:g2398683; PIDN:CAB16178.1; PID:g2398713 Smith, D.R.; Robison, K.
                                                                                                                                                                                                                                     Molecule type: DNA
Residues: 1-311 <PAR>
                                                                                                                                                                                                                                                                                                                                                Reference number: Z16918
                                                                                                                                                                                                                                                                                                                                                                                                                          Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 17-Nov-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Species: Mycobacterium leprae
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Best Local Similarity
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                                                                                                                                                                                                                                                                                            Status:
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Experimental source: strain 168
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Residues: 1-266 <K
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190, 249-256, 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Accession: F69974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Authors: Lauber, J.; Lazarevic, V.; Lee, S.M.; Levine, A.; Liu, H.; N.
M.; Ogawa, K.; Ogiwara, A.; Oudega, B.; Park, S.H.; Parro, V.; Pohl.
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S.; Brouillet, S.; Bruschi, C.V.; Caldwell, B.; Capuano, V.; Carter, N.M.; Chc
ch, S.D.; Emmerson, P.T.; Entian, K.D.; Errington, J.; Fabret, C.; Ferrari, E.
preliminary
e type: DNA
s: 'M',90-311 <SMI>
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33.3%;
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36.4%;
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Pred. No. 23;
6; Mismatches
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Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S.M.; Levine, A.; Liu, H.; Masuda,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B.; Rose, M.; Sadaie,
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7.5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lapidus, A.; Lardinois
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A;Cross-references: EMBL:U00013; NID:g466868; PIDN:AAA17144.1; PID:g466899 C;Genetics:
A;Gene: MICL536.34
A;Start codon: GTG
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Query Match 55.9%; Score 38; DB 2; Length 311; Best Local Similarity 58.3%; Pred. No. 27; Matches 7; Conservative 1; Mismatches 4; Indels 0; Gaps

Qy 1 VWFHVLFYLAVG 12 || || : || Db 151 VWLAVLLYVKVG 162

Search completed: June 4, 2001, 12:15:47 Job time: 280 sec

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
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## ALIGNMENTS

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	an email to license@isb-sib.ch).	requires a license agreement (See http://www.isb-sib.ch	ified and this statement is not removed. Usage by and for commerc	by non-profit institutions as long as its content is in no	the European Bioinformatics Institute. There are no restrictions on its	Detween the Swiss Institute of Bioinformatics and the EMBL outstation -	This SWISS-PROT entry is copyright. It is produced through a collaboration			<pre>-!- DATABASE: NAME=PROW; NOTE=CD guide CD64 entry;</pre>		STALL START - UNITED THE CONTRACT OF THE CONTR	IFICITY: MONO	ALTERNATIVE SPLICING.	PRODUCTS:	-!- SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN.	AFFINITY RECEPTOR.	ence 243:378-381(1989).	Complementary DNAs.";	isolation and expression or functional night-affinity to receptor	יה היים ליים ליים ליים ליים ליים ליים לי	1	MEDI THE ROLL OF THE MEDICAL PROPERTY OF THE P	SHOULD PROM N A	Nucleic Acids Res. 16:11824-11824(1988).	"Nucleotide sequence of three cDNAs for the human high affinity Fc	Allen J.M., Seed B.;	MEDLINE=89098339; PubMed=2974947;	SEQUENCE FROM N.A.		NCBI_TaxID=9606;	Primates; Catarrhini; Hominidae;	hordata;	HOmo sapiens (Human).	FCGR1A OR FCGR1 OR FCG1 OR IGFR1.	RI) (FCRI) (IGG FC RECEPTOR I) (CD64).	TY IMMUNOGLOBUL	(Rel. 40, Last	(Rel. 12,	01-OCT-1989 (Rel. 12, Created)	2315;.	FCG1 HUMAN STANDARD: PRT: 374 AA.	HUMAN	

PIR; PIR;

EMBL; X14356; CAA32537.1; -. EMBL; X14355; CAA32536.1; -.

PIR;

; S03018; S03018. ; S03019; S03019. ; A41357; A41357. ; B41357; B41357.

InterPro; IPR003006;
Pfam; PF00047; ig; 3.

146760; -.

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Best Local S
Matches 12
                                   Osman N., Kozak C.A., Macketter and mapping of the gene gamma RI and chromosomal location of the gamma RI and chromosomal for the gene J. Immunol. 148:1570-1575(1992)
                                                             SEQUENCE FROM N.A.
MEDLINE=92166399; PubMed=1531670;
Osman N., Kozak C.A., McKenzie I.F.,
          FCG1_MOUSE
P26151;
01-MAY-1992
                                                                                                                                                                                                                                  LT 2
_MOUSE
                                                                                                  MEDLINE-90111035; PubMed-2136886;
Sears D.W., Osman N., Tate B., McK
"Molecular cloning and expression
receptor for IgG.";
                                                                                                                                                                      01-MAY-1992 (Rel. 22) Last sequence update)
15-JUL-1999 (Rel. 38, Last amoutation updat
HIGH AFFINITY IMMUNOGLOBULIN GAMMA FC RECEPTOR I).
RI) (FCRI) (IGG FC RECEPTOR I).
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Mammalia; Eutheria; Rodentia;
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IG-LIKE C2-TYPE DOMAIN.
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P31436;
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MEDLINE-99226230; PubMed-10209755
Liu J.Y., Miller P.F., Gosink M.,
"The identification of a new fami
                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN-K12 / MG1655;
MEDLINE-93315143; PubMed=7686882;
MEDLINE-93315143; PubMed=7686882;
Burland V.D., Plinkett G. III, Daniels D.L., Blattner F.R.;
"DNA sequence and analysis of 136 kilobases of the Escherichia genome: organizational symmetry around the origin of replicatio Genomics 16:551-561(1993).
                                                                                                                                                                                                                                                                                                                                                                                                              Escherichia coli.
Bacteria; Proteobacteria;
Escherichia.
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entities requires a license
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A46480; A46480.
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ean Bioinformatics Institute.
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X TRANSPORTER
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31:1845-1851(1999)
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"T "Analysis of the Escherichia coli genome.
T region from 87.2 to 89.2 minutes.";
Nucleic Acids Res. 21:3391-3398(1993).
-! SUBCELLULAR LOCATION: INTEGRAL MEMBRANE
-!- SIMILARITY: TO E. COLI YQCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHARACTERIZATION.
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           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                           Caenorhabditis elegans by comparative Genome Res. 10:703-713(2000).
                                                                                                         SEQUENCE FROM N.A
MEDLINE-20272150;
                                                                                                                                                                                                                                                            TISSUE=Adrenal gland;
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                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                Lai C.-H., Chou C.-Y.,
                                                                                                                                                                         "Gene
                                                                                                                                                                                                                                               MEDLINE-20402571; PubMed-10931946;
                                                                                                                                                                                                                                                                                             NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                              -!- SIMILARITY: BELONGS TO THE UPF0101
                                                                                 MEDLINE=20272150; PubMed=10810093;
Lai C.-H., Chou C.-Y., Ch'ang L.-Y.,
"Identification of novel human genes
                                                                                                                                             IR.-M., Han Z.-G., Song H.-D., Peng Y.-D., Huang Q.-H., Ren S.-X., I Y.-J., Huang C.-H., Li Y.-B., Jiang C.-L., Fu G., Zhang Q.-H., B.-W., Dai M., Mao Y.-F., Gao G.-F., Rong R., Ye M., Zhou J., S.-H., Gu J., Shi J.-X., Jin W.-R., Zhang C.-K., Wu T.-M., ang G.-Y., Chen Z., Chen M.-D., Chen J.-L., lene expression profiling in the human hypothalamus-pituitary-adrenal its and full-length cDNA cloning.";
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el. 39, Last sequence update)
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Microbiology 143:2939-2943(1997).
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"Sequence of the Bacillus subtilis genome
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                                                                                                                                                                                                                                                                                                                                                                                            Bacillus subtilis. ~
Bacteria; Firmicutes; Ba
Bacillus/Staphylococcus
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SIMILARITY: BELONGS TO THE FNT FAMILY OF TRANSPORTERS (TC 2.44).
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an email to license@isb-sib.ch)
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Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomyces.
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Science 272:1353-1356(1996).
-!- CATALYTIC ACTIVITY: ACYL-COA
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SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.
                                                                                                                                                                                                                                           S19461; S19461.
S0000644; ARE1.
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                                                                                                                                                                                                                                                                                                                                                   requires a license
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                                                                                                                                                                                                                         IPR002688;
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                                                                                                                                                                               Transmembrane;
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Rothstein
        W.
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he EMBL/GenBank/DDBJ databases
                                                                                                                                                      Endoplasmic reticulum;
POTENTIAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A., Deckelbaum R.J., Sturley S.L.;
                                                                                                                                                                                                                                                                                                                                                                                  There are no rest
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Query Match Best Local S Matches 6

Similarity 54. 6; Conservative

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Score 38; DB Pred. No. 46; 2; Mismatches

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                                                                                                                                                                                                                                                                                                                                                   TRANSMEM SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kaneko T., Sato S., Kotani H., Tanaka A., Asamizu E., Nakamura Y., Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T., Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T., Hosouchi T., Matsuno A., Muraki A., Nakazaki N., Naruo K., Okumura Shimpo S., Takeuchi C., Wada T., Watanabe A., Yamada M., Yasuda M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synechocystis sp. (strain PCC 6803).
Bacteria; Cyanobacteria; Chroococcales;
NCBI_TaxID=1148;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Sequence analysis of the genome of the unicellular cyanobacterium Synechocystis sp. strain PCC6803. II. Sequence determination of the entire genome and assignment of potential protein-coding regions.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. MEDLINE=97061201; PubMed=8905231;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      YH90_SYNY3
P72793;
                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                             or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
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NCBI_TaxID=10116;
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                                                                                                                                                                                                                                                                                                                                                                                                   Hypothetical protein;
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by non-profit institutions as long as its content
ified and this statement is not removed. Usage by an
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SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (POTENTIAL).
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                                                                                                                                                                                                                                                                                 Similarity 77.
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24063 MW;
                         Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae; Rat
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77.8%;
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POTENTIAL.
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                          Murinae; Rattus
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VARSPLIC
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"Rat CD16 is defined by a family of class III Fc grequiring co-expression of heteroprotein subunits.
Inmunol. 146:4352-4361(1991).
                                                                                                                                                                                                                                                                             EMBL;
                                                                                                                                                                                                                                                                                                         EMBL;
                                                                                                                                                                                                                                                                                                                                                                            use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GAMMA. LOW AFFINITY RECEPTOR.

-!- SUBUNIT: MAY FORM MULTISUBUNIT COMPLEX WITH OTHER HETEROPROTEINS

-!- SUBUNIT: MAY FORM MULTISUBUNIT COMPLEX WITH OTHER HETEROPROTEINS
 VARIANT
                                                                                                                                                                                                                                 InterPro; IPR003006; -. Pfam; PF00047; 1g; 3.
                                                                                                                                                                                                                                                                                                                                                                or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Rat class III Fc gamma receptor isoforms binding specificity and fail to associate
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SEQUENCE FROM N.A. (1502-0-)
SEQUENCE FROM N.A. (1502-0-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-SPRAGUE-DAWLEY; TISSUE-Brain; MEDLINE-93246650; PubMed-8482840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Characterization and expression of an from rat natural killer cells.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=90239026; PubMed=1692135;
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                                                                                                     DOMAIN
                                                                                                                                                                                                      IgG-binding protein;
Immunoglobulin domai
                                                                                                                                                                                                                                                                HSSP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DIFFERENTIALLY REGULATED. TISSUE SPECIFICITY: EXPRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: CONTAINS 2 IMMUNOGLOBULIN-LIKE C2-TYPE DOMAINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EXPRESSION. DOES NOT ASSOCIATE WITH CD3 ZETA. SUBCELLULAR LOCATION: TYPE I MEMBRANE PROTEIN (POTENTIAL)
                                                                                                                                                                                                                                                             M32062; AAA41148.1;
L08446; AAA41151.1;
P12319; lALT.
                                                                                                                                                                                                                                                                                                      M64368; AAA42049.1;
M64369; AAA42048.1;
M64370; AAA42050.1;
                                                                                                                                                                                                                                                                                                                                                                 an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sci. U.S.A. 87:3425-3429(1990).
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241
267
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 39
                                                                                                                                                                                                                                                                                                                                                                             license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                     Receptor;
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IG-LIKE C2-TYPE DOMAIN.
IG-LIKE C2-TYPE DOMAIN.
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BY SIMILARITY.
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                                                                                                                                                                          LOW AFFINITY IMMUNOGLOBULIN
                                                                                                                                                                                                                   Transmembrane;
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productively with rat C
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Matches '6
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                                                                                                                                                                                                                                         MEDLINE-99254548; PubMed-10321245;
Patel A.J., Honore E., Lesage F., Fink M., Romey G., Lazdunski M.,
"Inhalational anesthetics activate two-pore-domain background K+
                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse).
Mus musculus (Mouse).
Morvota; Metazoa; Chordata;
                                                                                                                                                                                             NAT. NEUROSCI. 2:422-426(1999).
1- FUNCTION: VOLTAGE INSENSITIVE, INSTANTANE
POTASSIUM CHANNEL. OUTWARD RECTIFICATION
EXTERNAL K+ CONCENTRATIONS.
                                                                                                                                                                                                                                                                                              "A neuronal two P domain K+ channel stimulated by arachidonic acid polyunsaturated fatty acids."; EMBO J. 17:3297.3308(1998).
                                                                                                                                                                                                                                                                                                                                       MEDLINE=98292450; PubMed=9628867;
Fink M., Lesage F., Duprat F., He
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SEQUENCE
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01-OCT-2000
                                                                                                                                                                                                                                                                                 ACTIVATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          POTASSIUM CHANNEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                217 VWFHAAFCLVM 227
                                                                          MISCELLANEOUS: ACTIVATED BY ARACHIDONIC ACID AND OTHER UNSATURAL FATTY ACIDS. NOT AFFECTED BY VOLATILE GENERAL ANAESTHETICS SUCH CHLOROFORM, DIETHYL ETHER, HALOTHANE AND ISOFLURANE. SIMILARITY: BELONGS TO THE TWO PORE DOMAIN FAMILY OF POTASSIUM
                                                                                                                                       SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (POTENTIAL). ALTERNATIVE PRODUCTS: 2 ISOFORMS; 1 (SHOWN HERE) AND 2/TRAAKT/TRUNCATED; ARE PRODUCED BY ALTERNATIVE SPLICING. TISSUE SPECIFICITY: EXPRESSED IN BRAIN, SPINAL CORD AND EYE.
                                                                                                                                                                                   SUBUNIT: HOMODIMER (POTENTIAL).
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                                                                                                                                  DETECTED IN HEART,
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(Rel. 40, Last annotation update)
ANNEL SUBFAMILY K MEMBER 4 (TWIK-RELATED ARACHIDONIC ACID-
OTASSIUM CHANNEL PROTEIN) (TRAAK).
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It is produced through a cinformatics and the EMBL

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the European Land the European Land the European Land this statement is not removed. modified and this statement is not removed. entities requires a license agreement (See and an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                    CRTI_CERNC
P48537;
01-FEB-1996
                                                                                                                                                       nicotianae phytoene dehydrogenase gene.";
Appl. Environ. Microbiol. 60:2766-2771((1994).
-i- FUNCTION: THIS ENZYME CONVERTS PHYTOENE INTO ZETA-CAROTENE VIA THE
INTERMEDIARY OF PHYTOFLUENE BY THE SYMMETRICAL INTRODUCTION OF TWO
DOUBLE BONDS AT THE C-11 AND C-11' POSITIONS OF PHYTOENE.
                                                                                                                                                                                                                                                                                                       Eukaryota; Fungi; Ascomycota; Pezizomycotina;
Dothideomycetes et Chaetothyriomycetes incert
Mycosphaerellaceae; mitosporic Mycosphaerella
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01-FEB-1996
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VARSPLIC
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Ehrenshaft M., Daub M.E.;
                                                                                                                                                                                                                                                                                                                                           Cercospora nicotianae.
                                                                                                                                                                                                                                                                                                                                                                  PHYTOENE DEHYDROGENASE
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                                   ween the Swiss Institute. Tree European Bioinformatics Institute. Tree mon-profit institutions as long
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                                                                                                       SIMILARITY: TO BACTERIAL PHYTOENE DEHYDROGENASES AND TO METHOXYNEUROSPORENE DEHYDROGENASE (CRTD).
                                                                                                                                             COFACTOR: FAD (PROBABLE).
                                                                                                                                  PATHWAY: CAROTENOID BIOSYNTHESIS.
                                                                                SWISS-PROT entry is copyright. It is produced through a collaboration
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Pred. No. 47;
1; Mismatches
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N-LINKED (GLCNAC. . .) (POTENT.
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KLLVE -> KAMAI (IN ISOFORM 2).
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P51944; Q60797;
01-OCT-1996 (Rel
                                                                                                                                                                                                                                                                          Croci L., Stayton C.L., Bossolasco M., Bianchi E., Corradi A.M., Pardi R., Consalez G.G.;
Submitted (MAR-1995) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: LIKELY TO BE INVOLVED IN THE CONTROL OF THE CELL CYC DURING S PHASE AND G2 (BY SIMILARITY).
-I- SUBÇELLULAR LOCATION: NUCLEAR (BY SIMILARITY).
-I- SUBÇELLULAR LOCATION: THE CYCLIN FAMILY. CYCLIN AB SUBFAMILY.
-I- SIMILARITY: BELONGS TO THE CYCLIN FAMILY.
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TRANSMEM
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01-OCT-1996 (Rel. 34, Last sequence up
01-OCT-2000 (Rel. 40, Last annotation
G2_MITOTIC-SPECIFIC CYCLIN F.
                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE-95284479; PubMed-7767003;
Obermayr F.O., Sutherland H.F., Kraus B., Frischauf A.-M.;
"Mouse cyclin F maps to a conserved linkage group on mouse chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00982; PHYTOENE_DH; 1.
Carotenoid biosynthesis; Oxidoreductase; FAD; Flavoprotein;
Cyclin; Cell cycle; Cell division; Mitosis; Nuclear protein; Alternative splicing.
                                  InterPro; IPR000553; -.
InterPro; IPR001810; -.
Pfam; PF00646; F-box; 1.
Pfam; PF00134; Cyclin; 1.
PROSITE; PS00292; CYCLINS; 1.
                                                                                                                                                                                       entities
                                                                                                                                                                                                                                          between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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                         PROSITE; PS50181; FBOX; 1.
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                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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; U20612; AAA62317.1;
; U20636; AAA63152.1;
; P30274; IVIN.
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                                                                                                                                                                           equires a license agreement (semail to license@isb-sib.ch)
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621 AA;
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Pallavicini A., Lanfranchi G., Valle G.;
"The DNA sequence of cosmid 14-13b from chromosome XIV of
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                                                                                                                                                                                                                                                                                                                   Saccharomyces cerevisiae reveals an unusually high number of overlapping open reading frames.";
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Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
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Transmembrane
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GALACTOSE-1-PHOSPHATE URIDYLYLTRANSFERASE (EC
GAL7 OR YBR018C OR YBR0226
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Nolling J. Reeve J.N.;
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"Growth- and substrate dependent transcription of
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-1- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN (PROBABLE)
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    (Baker's yeast).
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POTENTIAL.
POTENTIAL.
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Pred.
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B0D047347BF7B1A4 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 update)
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                                                     2.7.7.10)
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"Sequence and functional analysis of a 7.2 kb fragment of Saccharomyces cerevisiae chromosome II including GAL7 and new essential open reading frame."; yeast 11:79-83(1995).
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Tajima M., Nogi Y., Fukasawa T.;
"Primary structure of the Saccharomyces cerevisiae GAL7 gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                  ACT_SITE
ACT_SITE
CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 1-184 FROM N.A. STRAIN=CARLSBERGENSIS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=S288C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Citron B.A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast 1:67-77(1985).
                                       CONFLICT
                                                                                                                                                                                                                                                       EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       yeast GAL7 gene.";
Nucleic Acids Res. 11:8555-856
-i- CATALYTIC ACTIVITY: UTP +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                             or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                                                                                                           entities requires a
                                                                                                                                                                                                                                                                                                                                      modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the El
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=84185433; PubMed=6715281;
                                                                                                                                                                                                                                                                                   EMBL;
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SEQUENCE
            CONFLICT
                          CONFLICT
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                                                                                                                                                 Pfam;
                                                                                                                                                                                                                  PIR; S45873; XNBYUG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nucleotide sequence of the
                                                                                                                                                              InterPro;
                                                                                                                                                                           InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacteriol. 158:269-278(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                       URIDYLYLTRANSFERASE FAMILY 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                     SIMILARITY: BELONGS
                                                                                                                                                                                                                                                                                                                                                                                                                                                PATHWAY: GALACTOSE METABOLISM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             UDP-GALACTOSE +
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M12348;
X00215;
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                                                                                                                                    01087; GalP_UDP_transf; 1.
PS00117; GAL_P_UDP_TRANSF_I;
                                                                                                                                                                                                                                                                                                                                                    non-profit institutions as long
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                                                                                                                         Nucleotidyltransferase;
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AAA34627.1;
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CAA84960.1;
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V -> A (IN REF. )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Schindwolf T., Lehnert W.,
                                                     H -> Y (IN |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALPHA-D-GALACTOSE 1-PHOSPHATE
                                                                                  PROBABLE
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  EBA67A8722D19F3B CRC64;
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            IN RE. (IN REF. 3).
S (IN REF. 1).
TN REF. 1).
                                                                                                                           Galactose metabolism
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Query Match

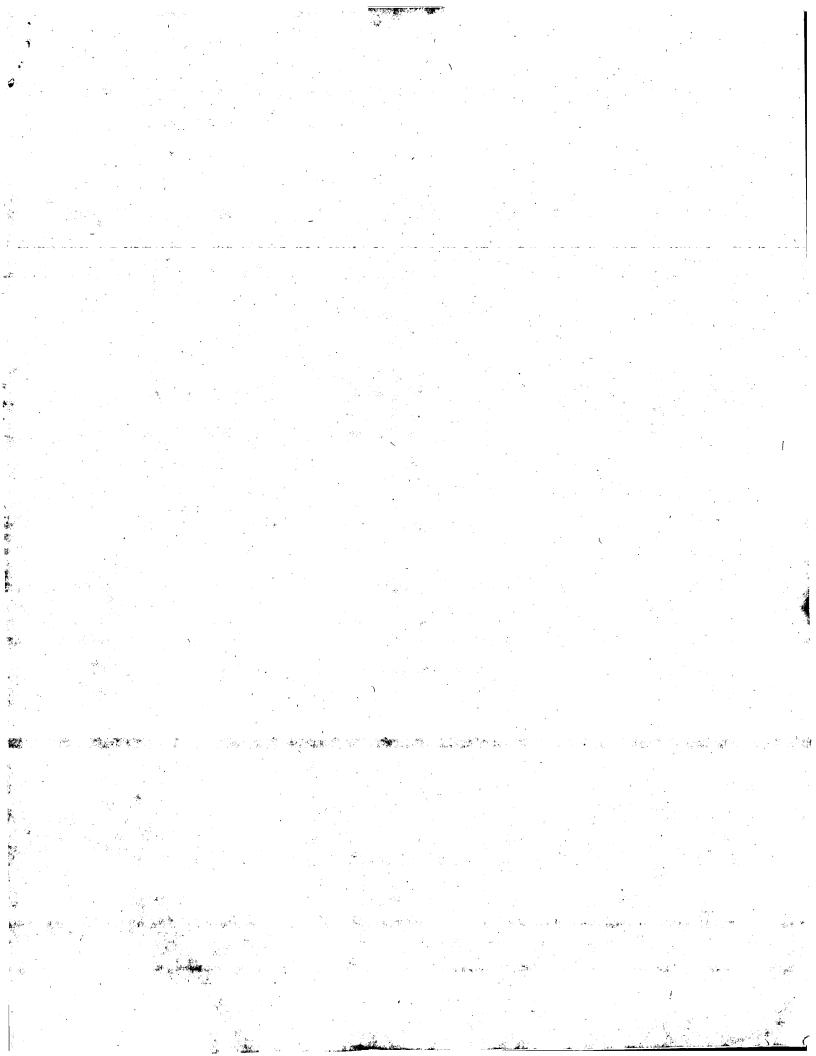
52.9%;

Score 36;

DВ

۲.

Length 365



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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein search, using sw model
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1: sp_archea:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 June 4, 2001, 12:23:31; Search time 107.68 Seconds (without alignments)
13.062 Million cell updates/sec
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Gapop 10.0 , Gapext 0.5
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68
1 VWFHVLFYLAVG 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                           sp_rodent:*
sp_unclassified:*
sp_vertebrate:*
sp_virus:*
                                                                                                                                                                                                                                                                                  sp_bacteria:*
sp_fungi:*
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## SUMMARIES

Result	Score	% Query Match	Length	DВ	ID.	• *	Description
1	68	100.0	188	4	Q92638	1	Q92638 homo sapien
N	68	100.0	280	4	Q92637		homo.
w	. 68	100.0	374	4	Q92663		homo
4	68	100.0	375	4	Q92495		homo
ъ	63	92.6	349	σ	Q9MZTO		Q9mzt0 bos taurus
σ	45	66.2	460	N	Q9xD16		Q9xd16 leptospira
7	.41	60.3	153	N	Q9WXU4		Q9wxu4 thermotoga
8	41	60.3	278	14	P89139		P89139 simian hemo
9	. 41	60.3	340	ഗ	018731		Q18731 caenorhabdi
10	41	60.3	1852	ω	013281		
11	41	60.3	1867	w	Q9P8W4		09p8w4 blumeria gr
12	40	58.8	88	14	Q89405		Q89405 paramecium
13	40	58.8	266	N	052498		O52498 bradyrhizob
14	40	58.8	448	Ŋ	031428		031428 bacillus su
15	39	57.4	168	Ņ	Q9RQ17		Q9rqi7 listeria mo
16	39	57.4	378	10	Q9M3W5		Q9m3w5 avena sativ
17	39	57.4	1075	10	Q9LNP5		Q9lnp5 arabidopsis
18	38.5	56.6	272	4	095004		095004 homo sapien
19	38.5	56.6	1084	4	Q9UPP1		

	45	.44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20
	37	37	37	37	37	37	37	37	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38	38
	54.4	54.4	54.4	54.4	54.4	54.4	54.4	54.4	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9	55.9
	194	190	148	148	148	148	148	148	1869	907	4.97	398	395	393	391	357	354	348	343	314	311	223	197	172	111	84
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	047579	Q9U5Q4	Q9T3U4	Q9T3V8	Q9T414	Q9TH29	Q9TH30	Q9TH31	093809	OMZA60	Q9W1D4	Q9SXX5	Q9SXX4	Q9SJU4	Q9ZU52	P93565	Q08417	Q9W280	Q9SVJ6	058219	033142	Q49704	Q9RKU2	Q9TTU2	Q9XEF7	026862
-	047579 onchocerca	Q9u5q4 entodinium	Q9t3u4 campeloma	Q9t3v8 campeloma	Q9t414 campeloma	Q9th29 campeloma	Q9th30 campeloma	. Q9th31 campeloma	O93809 magnaporthe	Q9vzw0 drosophi	Q9w1d4 drosophi	Q9sxx5 nicotiana	Q9sxx4 nicotiana		Q9zu52 arabi	P93565 solan	. Q08417 saccharomyc	Q9w280 drosophila	Q9svj6 arabidopsis	058219 pyrococcus	<ul> <li>O33142 mycobacter</li> </ul>	Q49704 mycobacter	Q9rku2 streptomyce	Q9ttu2 oryctolagus	Q9xef7 arabidopsis	. 026862 methanobact
	erca	nium	oma 1	oma 1	oma p	oma l	oma p	oma p	orthe	hila	hila	iana p	iana p	arabidopsis	arabidopsis	solanum tub	romyc	hila	dopsis	ccus	cteri	cteri	отусе	lagus	dopsis	obact

## ALIGNMENTS

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л 2 7 Q92637 PRELIMINARY; PRT; 280 AA.	1 VWEHVLEYLAVG 12             103 VWEHVLEYLAVG 114,	Query Match 100.0%; Score 68; DB 4; Length 188; Best Local Similarity 100.0%; Pred. No. 0.00025; Matches 12; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	SEQUENCE 188 AA; 22106 MW; E81517B33BEAB789 CRC64;	PRODOM; PD002534; -; 1.	PEAM; PF00047; ig; 1.	Ē.	EMBL; L03420; AAA35826.1; -	in. Inve	1	"Novel Fc gamma receptor I family gene products in human mononuclear	ly R.P.;	$\sim$	MEDLINE=93055454; PubMed=1430234;		SEQUENCE FROM N.A.			Primates; Catarrhini; Hominidae;	ryota; Me	Homo sapiens (Human).	B2.	FC GAMMA RECEPTOR I	(TrEMBLrel. 15, Last annotation	02,	01-FEB-1997 (TrEMBLrel. 02, Created)	Q92638;	Q92638 PRELIMINARY; PRT; 188 AA.	00	9 P	
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01-FEB-1997 (TremB)
01-FEB-1997 (TremB)
01-QCT-2000 (TremB)
FC GAMMA RECEPTOR
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Q92663;
01-FEB-1997
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=93055454;
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                                                                                                                                                                     INTERPRO; IPR003006; -. PFAM; PF00047; ig; 3. PRODOM; PD002534; -; 1. SEQUENCE 374 AA; 426
                                                                                                                                                                                                                                                 EMBL; L03418; AAA36049.1;
HSSP; P12319; 1ALT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-FEB-1997 (TrEMBLrel. 01-OCT-2000 (TrEMBLrel.
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Kimberly R.P.;
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                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=93055454;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9606;
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                                                                                             Local Similarity
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                                     1 VWFHVLFYLAVG
                                                                                                                                                                                                                                                                                       Clin. Invest. 90:2102-2109(1992)
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P12319; 1ALT.
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Primates;
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Primates;
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32232 MW;
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                                                                       Score 68; DB 4; red. No. 0.00048;
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Pred. No.
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Ц.Е.
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01-FEB-1997 (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Definition of interferon gamma-response elements in a novel human Fc gamma receptor gene (Fc gamma RIb) and characterization of the gene structure ".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Benech P.D., Sa
Ezekowitz A.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
Bovidae; Bovinae; B
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               SEQUENCE FROM N.A.
                                                                                                                                           Bos taurus (Bovine)
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                                                                                Bos.
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Primates;
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                                                                                                    Chordața; Craniata;
Cetartiodactyla; Ru
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Pred. No. 0.00048;
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                                                                                                      Ruminantia; Pecora;
                                                                                                                                                                                                                                                                          349
                                                                                                        Vertebrata; Euteleostomi;
minantia; Pecora; Bovoidea;
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zuerner R.L., Harskeer R.A., van de Kemp H., Bal A.E.;

*Sequence analysis of a large ribosomal protein gene operon
Leptospira interrogans.";

Submitted (DEC-1998) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: INVOLVED IN PROTEIN EXPORT.

-!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.

-!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN.

-!- SIMILARITY: BELONGS TO THE SECY/SEC61-ALPHA FAMILY.

EMBL; AF115283; AAD40603.1;
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STRAIR-SEROVAR LAI;
MEDLINE-94041964; PubMed-8409911;
Gravekamp C., van de Kemp H., Franzen M., Carrington D.,
Van Eys G.J., Everard C.O., Hartskeerl R.A., Terpstra W.J
"Detection of seven species of pathogenic leptospires by
sets of primers.";
J. Gen. Microbiol. 139:0-0(0).
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EMBL; AF162866; AAF80477.1; ..
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                                                                                                                                                                                                                                                                  PRINTS; PR00303; SECYTRNICASE.
PROSITE; PS00018; EF_HAND; UNKNOWN_1.
PROSITE; PS00095; C5_MTASE_2; UNKNOWN_1.
PROSITE; PS00755; SECY_1; 1.
PROSITE; PS00756; SECY_2; 1.
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"Evidence for lateral gene transfer between Archaea and Bacteria from genome sequence of Thermotoga maritima.";

Nature 399:323-329(1999).

EMBL; AE001695; AAD35183.1; -.
                SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=LVR 42-0/M6941;
MEDLINE=97361760; PubMed=9218721;
MEDLINE=97361760; PubMed=97361760;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Viruses; ssrNA positive-strand NCBI_TaxID=38143;
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STRAIN-MSB8 / DSM 3109;
MEDLINE-99287316; PubMed-10360571;
Nelson K.E., Clayton R.A., Gill S.R.,
Haft D.H., Hickey E.K., Peterson J.D.,
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  Gene
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J. Virol. 69:2679-2683(1995).
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191:205-210(1997)
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SEQUENCE FROM N.A. STRAIN-LVR 42-0/M6941;

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Smith

S.L.,

Wang

X.C.,

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013281 PRELIMINARY;
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PRODOM; PD001151; -; 1.
SEQUENCE 278 AA; 31294 MW;
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Bonfield J., Burton J., Connell M., Copsey T., Coulson A
Craxton M., Dear S., Du Z., Durbin R., Favello A., Fulton L.,
Gardner A., Green P., Hawkins T., Hillier L., Jier M., Johnston L.,
Jones M., Kershaw J., Kirsten J., Laister N., Latreille P.,
Lightning J., Lloyd C., Mcmurray A., Mortimore B., O'Callaghan M.,
Parsons J., Percy C., Rifken L., Roopra A., Saunders D., Shownkeen
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J.,
Smaldon N., Smith A., Sonnhammer E., Staden R., Sulston J.
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Nature 368:32-38(1994).

EMBL. Z72505; CAA96610.1; -. F9E23EA24761F151 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditidae; Peloderinae; Caenorhabditis. NCBI_TaxID-6239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C50C10.4 PROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Caenorhabditis elegans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C50C10.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     IEDLINE=94150718; PubMed=7906398;
                                                                                                                                    282
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                                                                                                                                   WYHVLFLRPIG 292
                                                                                                                                                           WEHVLFYLAVG
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                                                                                                                                                                                      Similarity
6; Conserv
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                                                                                                                                                                                    Score 41; DB Pred. No. 20; 2; Mismatches
Last sequence update)
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.16) (CHITIN-UDP ACETYL-GLUCOSAMINYL
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                                         Created)
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                                                                  1852 AA
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Fungal Genet. Biol. 20:153-167(1996).

Fungal Genet. Biol. 20:153-167(1996).

FUNCTION: CHITIN SYNTHESIZED BY CHITIN SYNTHASE D CONTRIBUTES TO FUNCTION: CHITIN SYNTHESIZED BY CHITIN GEONIDIA, OF THE SUBAPICAL THE RIGIDITY OF THE WALLS OF GERMINATING CONIDIA, OF THE SUBAPICAL REGION OF HYPHAE, AND OF CONIDIOPHORE VESICLES, BUT IS NOT NECESSARY FOR NORMAL MORPHOLOGY OF THESE CELLS.

-i-- CATALYTIC ACTIVITY: UDP-N-ACETYL-D-GLUCOSAMINE + [1,4-(N-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-ACETYL-IN-
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Winter K.R., Riggle P.J.,
Borgia P.T.;
                                                                                                                                                                                                                                     VARIANT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TRANSMEM TRANSMEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE OF 359-1852 FROM N.A. STRAIN=FGSC 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. STRAIN=FGSC A89;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Fungi;
NCBI_TaxID=5072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Emericella nidulans (Aspergillus nidulans).
Eukaryota; Fungi; Ascomycota; Eurotiales; Trichocomaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CSMA OR CHSD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS01039; SBP_BACTERIAL_3; PRODOM; PD000355; -; 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- SIMILARITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- SUBCELLULAR LOCATION: INTEGRAL MEMBRANE PROTEIN. PLASMA MI
-!- DOMAIN: CONTAINS A MYOSIN-LIKE GLOBULAR HEAD DOMAIN FUSED
CHITIN SYNTHASE C-TERMINAL DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "The chsD and chsE genes of Aspergillus nidulans and their roles
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PFAM; PF00063; myosin_head;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INTERPRO; IPR001638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INTERPRO; IPRO01173; INTERPRO; IPRO01199;
                                                                                                                                                                                                                                                                                                                                                                         FRANSMEM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENTERPRO;
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3 FHYLFYLAVG 12
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U62895; AAB05797.1;
                                                                  Similarity 70. 7; Conservative
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1852
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70.0%;
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P.J., Rhodes J.
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                                                                     1;
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                                                                                                    Score 41;
Pred. No.
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MISSING (IN STAIN FGSC 4).

ATEAVLSGNL -> GKHRPI (IN S
ATEAVLSGNL -> DBD2D57AOD42617F CRC64
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POTENTIAL.
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                                                                     Mismatches
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.C., Dodge C.
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                                                                                             DB
99;
                                                                                                                                           ω,
                                                                                                                                           Length 1852;
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L., Culp D.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cell wall;
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CRC64;
                                                                            Indels
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Best Local Similarity 70.0
"""" 7; Conservative
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Q89405;
052498
052498;
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Eukaryota; Fungi;
NCBI_TaxID=34373;
                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE-95133167; PubMed-7831789;
MEDLINE-Y., Zhang Y., Kutish G.F
"Analysis of 45 kb of DNA located
                                                                                                                                                                                                                                                                                                                                                                                                                                        Zhang Z.G., Gurr S.J.,
Submitted (APR-2000) t
EMBL; AF189366; AAF042
SEQUENCE 1867 AA; 2
                                                                                                                                  Hypothetical protein. SEQUENCE 88 AA; 10265 MW;
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01-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 graminis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-CC1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CHITIN SYNTHASE.
                                                                                                                                                      EMBL;
                                                                                                                                                             Virology
                                                                                                                                                                                                                                    Paramecium bursaria chlorella virus 1
Viruses; dsDNA viruses, no RNA stage;
                                                                                                                                                                                                                                                                HYPOTHETICAL
                                                                                                                                                                                                                                                                                           01-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=CC1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2000
01-OCT-2000
                                                                                                                                                                                                                           NCBI_TaxID=10506;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Differential expression
                                                                                                                                                                                                                                                                                                                                                         196 FHVLYYLLAG
                                                    21
                                                                                      Local Similarity
nes 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                3 FHVLFYLAVG
                                                                   1 VWFHVLFYL 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plant Pathol.
                                                    IMFHILFYL
                                                                                                                                                   PBCV-1 genome.";
ogy 206:339-352(1995).
U42580; AAC96438.1; -.
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O (TrEMBLrel. 15,
O (TrEMBLrel. 15,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hall A.,
                                                                                       Conservative
                                                                                                                                                                                                                                                                       (TrEMBLrel. 01, (TrEMBLrel. 01, (TrEMBLrel. 08,
        PRELIMINARY;
                                                                                                                                                                                                                                                                                                            PRELIMINARY;
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                                                                                              58.8%;
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70.0%;
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sion of t
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two chitin synthase
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Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 41;
Pred. No.
                                                                                               Pred.
                                                                                               Score 40;
Pred. No.
                                                                                                                                   10382DA045764EED CRC64;
        PRT;
                                                                                                                                                                                                                                                                                                            PRT;
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                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                              Rock D.L., Van Etten J.L.; the left end of the chlorella
        266 · AA
                                                                                                                                                                                                                                    (PBCV-1).
Phycodnaviridae;
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1e+02;
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7.9;
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                                                                                                      Length 88
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Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      O31428;
O1-JAN-1998
O1-JAN-1998
O1-MAY-2000
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01-JUN-1998
01-JUN-2000
                                                                                                                                                                                                                                                                                                                     Borriss R., Boursier L., Brans A., Braun M., Brignell S.C., Bron S., Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M., Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A., Denizot F., Devine K.M., Dusterhoft A., Ehrlich S.D., Emmerson P.T., Entian K.D., Errington J., Rabret C., Ferrari E., Foulger D., Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N., Ghim S.Y., Glaser P., Goffeau A., Golightly E.J., Grandi G., Guiseppi G., Guy B.J., Haga K., Haiech J., Harwood C.R., Henaut A., Guiseppi G., Guy B.J., Haga K., Haiech J., Harwood C.R., Henaut A., Jones L., Holsappel S., Hosono S., Hullo M.F., Itaya M., Jones L., Joris B., Karamata D., Kasahara Y., Klaerr-Blanchard M., Klein C., Kubayashi Y., Koetter P., Koningstein G., Krogh S., Kumano M., Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V., Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V., Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V., La
                                                                                                              Kurita K., Lapidus A., Lardinois S., Lauber J., Lazarevic V.,
Lee S.M., Levine A., Liu H., Masuda S., Mauel C., Medigue C.,
Medina N., Mellado R.P., Mizuno M., Moestl D., Nakai S., Noback M.
Noone D., O'Reilly M., Ogawa K.; Ogiwara A., Oudega B., Park S.H.,
Parro V., Pohl T.M., Portetelle D., Porwollik S., Prescott A.M.,
Presecan E., Pujic P., Purnelle B., Rappoplik S., Reynolds :
Rieger M., Rivolta C., Rocha E., Roche B., Rose M., Sadaie Y.,
Sato T., Scanlan E., Schleich S., Schroeter R., Scoffone F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Becker B.U., Kosch K., Parniske M., Muller r.; Submitted (DEC-1997) to the EMBL/GenBank/DDBJ databases. EMBL; AF039306; AAC04824.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bradyrhizobium japonicum.
Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacillus subtilis.
Bacteria; Firmicutes; Ba
Bacillus/Staphylococcus
NCBI_TaxID=1423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           031428
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Sekiguchi J., Sekowska A., Seror S.J., Serror P., Shin B.S., Sorokin A., Tacconi E., Takagi T., Takahashi H., Takemaru K. Takeuchi M., Tamakoshi A., Tanaka T., Terpstra P., Tognoni F. Tosato V., Uchiyama S., Vandenbol M., Vannier F., Vassarotti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kunst F., Ogasawara N., Moszer I., Albertini A.M., Alloni G., Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN-USDA 110SPC4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=98044033;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INTERPRO; IPR000051; INTERPRO; IPR001601;
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B (TrEMBLrel.
CTrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (TrEMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PubMed=9384377;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ria; alpha subdivision;
Bradyrhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13,
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14,
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Last sequence update)
Last annotation updat
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Last sequence up
Last annotation
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0318A45D2A1B958C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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23;
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                                                                                                                                                                             Rey M., Reynolds S.
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      Vassarotti A.,
                                                                                              Shin B.S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Borchert S.
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                                                                                           Soldo
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                                                                                                       Query Match
Best Local Similarity
Matches 5; Conserv
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Best Local Similarity 60.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPROI7 PRELIMINARY; PRT; 168 AA.
OPROI7;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-JUN-2000 (TrEMBLrel. 14, Last annotation update)
HYPOTHETICAL 18.7 KDA PROTEIN (FRAGMENT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-168;
Haga K., Liu H., Yasumoto K., Takahashi H., Yoshikawa H.;
Haga K., Liu H., Yasumoto K., Takahashi H., Yoshikawa H.;
"Sequence analysis of the 70kb region between 17 and 23 degree of Bacillus subtilis chromosome.";
Submitted (JUL-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; 299105; CAB11990.1;
EMBL; 299105; CAB11990.1;
INTERPRO; IPR000179;
PROSITE; PS00193; CYTOCHROME B_00; UNKNOWN_1.
SEQUENCE 448 AA; 51880 MW; 3870F3A47E445548 CRC64;
                                                                                                                                                                                                                                                                                                                                                                STRAIN-EGD;
FSihi H., Cossart P.;
FSihi H., Cossart P.;
"STM applied to Listeria monocytogenes.";
Submitted (OCT-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF102170; AAF04741.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN-168;
Kunst F., Ogasawara N., Yoshikawa H., Danchin A.;
Submitted (NOV-1997) to the EMBL/Genbank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Listeria monocytogenes.
Bactleria; Firmicutes; Bacillus/Clostridium group;
Bacillus/Staphylococcus group; Listeria.
NCBI_TaxID=1639;
                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nature 390:249-256(1997).
                                                                                                                                                                                                                                                                                           INTERPRO; IPR000566; -- ROSITE; PS00213; LIPOCALIN; UNKNOWN_1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EQUENCE FROM N.A.
86 IWFPVMAFVAIG 97
                         1 VWFHVLFYLAVG 12 :|| |: ::|:|
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                                                                                                                                                                                                                         168 AA; 18718 MW;
                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  58.8%; Score 40; DB 2; Length 448; 60.0%; Pred. No. 38; Five 3; Mismatches 1; Indels
                                                                                                                                Score 39; DB 2;
Pred. No. 22;
                                                                                                                                                                                                                                          9CF3891496C0EB8D CRC64;
                                                                                                            Mismatches
                                                                                                                                                             Length 168
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completed: June me: 595 sec

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